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Title:
Perfect score:
Sequence:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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233
1 gtttttgcaqttad
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| (Ggn2_6/ptodata/1/pubpna/USO7_PUBCOMB.seq:*
| (Ggn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
| (Ggn2_6/ptodata/1/pubpna/USO6_NEW_PUB.seq:*
| (Ggn2_6/ptodata/1/pubpna/USO6_PUBCOMB.seq:*
| (Ggn2_6/ptodata/1/pubpna/USO7_NEW_PUB.seq:*
| (Ggn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
| (Ggn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
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                                                                                                                                                  /cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US10C_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US10D_PUBCOMB.seq:*
cgn2_6/ptodata/1/pubpna/US10F_PUBCOMB.seq:*
cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
cgn2_6/ptodata/1/pubpna/US11_NEW_PUB.seq:*
cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
                                                                                                                                                                                                                                                                                                                                                                6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
6/ptodata/1/pubpna/US09A_PUBCOMB.seq:*
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                                                                                                                'ptodata/1/pubpna/US10C_PUBCOMB.seq:*
/ptodata/1/pubpna/US10D_PUBCOMB.seq:*
/ptodata/1/pubpna/US10E_PUBCOMB.seq:*
/ptodata/1/pubpna/US10E_PUBCOMB.seq:*
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5365.716 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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113.4	113.4	131.6	138	138	141	164.8	164.8	208.8	221.8	233	Score
48.7	48.7	56.5	59.2	59.2	60.5	70.7	70.7	89.6	95.2	100.0	Query Match
2583	1900	1924	1929	713	749	5580	1693	1770	2311	233	Query Match Length DB
18	100	18	18	18	18	18	18	.18	18	10	DB
US-10-437-963-508	115-10-437-963-509	US-10-739-930-5153	US-10-425-115-48185	US-10-767-701-12821	US-10-425-115-174626	US-10-437-963-91761	US-10-437-963-23894	US-10-425-115-160357	US-10-425-115-160360	US-09-237-183A-1252	ID
-	509	Sequence 5153, A	Sequence 48185, A	Sequence 12821, A	Sequence 174626,	Sequence 91761, 1	Sequence 23894, .	Sequence 160357,	Sequence 160360,	Sequence 1252, Ap	Description

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15 10 13	18 18 18 13	18 18 18	17 10 17 14 15 13	17 18 18	17 17 18 17
US-10-424-597-7034 US-10-311-455-556 US-09-918-995-29025 US-10-027-632-40743 US-10-027-632-40743	US-10-781-362-19 US-10-425-115-172445 US-10-674-124A-23429 US-10-027-632-139077 US-10-027-632-139077	US-10-781-362-13 US-10-781-362-9 US-10-781-362-9 US-10-781-362-5 US-10-781-362-7 US-10-781-362-17 US-10-781-362-17	-10-424-59 -09-844-653 -10-282-122 -10-239-676 -10-240-453 -10-027-632	US-10-425-114-5220 US-10-767-701-10807 US-10-437-963-83072 US-10-425-115-60567	-10
Sequence Sequence Sequence Sequence		sequence sequence sequence sequence sequence sequence sequence			
556, App 556, App 29025, A 40743, A	19, Appl 172445, 23429, A 239077, 139077,	13, Appi 9, Appli 11, Appl 5, Appli 7, Appli 15, Appli 17, Appl	83608, A 5, Appli 34479, A 13, Appl 21, Appl 21, Appl 139078, 139078,	· · · ›	, , , , , , , ,

ALIGNMENTS

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; TYPE: DNA
; ORGANISM: Zea mays
US-09-237-183A-1252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-237-183A-1252
                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Chisher, No. US20030135870Aldine
APPLICANT: Fisher, Dane K.
APPLICANT: Liu, Jingdong
TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With The TITLE OF INVENTION: Sucrose Pathway
FILE REFERENCE: 38-21(15089)B
CURRENT APPLICATION NUMBER: US/09/237,183A
CURRENT APPLICATION NUMBER: US/09/237,183A
CURRENT FILING DATE: 1999-01-26
PRIOR APPLICATION NUMBER: US 60/067,000
PRIOR APPLICATION NUMBER: US 60/067,000
PRIOR FILING DATE: 1997-11-24
NUMBER OF SEQ ID NOS: 2814
SEQ ID NO 1252
LENGTH: 233
                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 233; Conserv
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13
                              61 ATTGAATGAGGCAATCTCTGAGTATGAGACTTCAGAAAACAATGACTGCGGGAAGCTACCG 120
                                                                                                                        GTTTTTGCAGTTAGTAGAATATGTTAGTGGCTCCTATGATAGGGTGGAAGGATTTGAGTT 60
ATTGAATGAGGCAATCTCTGAGTATGAGACTTCAGAAAACAATGACTCGGGAAGCTACCG
                                                                                                                                                                                             100.0%; Score 233; DB 10; ilarity 100.0%; Pred. No. 6.4e-66; Conservative 0; Mismatches 0;
                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                           Length 233;
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APPLICANT: La ROSA, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules an
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
, SEQ ID NO 160357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Kovalic, David K.

APPLICANT: Kovalic, David K.

APPLICANT: Edou, Yihua

APPLICANT: Zhou, Yihua

APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Nucleic Acid Molecules and Other Title OF INVENTION: Plants

FILE REFERENCE: 38-21(5322)B

CURRENT APPLICATION NUMBER: US/10/425,115

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 369326

SEQ ID NO 160360

LENGTH: 2311

TYPE: DNA
ORGANISM: Zea mays
                                                                                                                                                                                                       RESULT 3

US-10-425-115-160357

; Sequence 160357, Application US/10425115

; Publication No. US20040214272A1

; GENERAL INFORMATION:
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Publication No. US20040214272A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
NAME/KEY: unsure
LOCATION: (1)..(2311)
OTHER INFORMATION: unsure at all n locations
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                           121 CAGATTATTTATTTGGCATTGCCTCCATCAGTCTACCCATCAGTATGCGAGATGATAAG 180
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97.0%;
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                                                                                                     and Other Molecules Associated With
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       182
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APPLICANT: Boukharov, Andrey A.

APPLICANT: Barbazuk, Brad

APPLICANT: Li, Ping

ITITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With

FILE REFERENCE: 38-21(53221)B

CURRENT APPLICATION UNMER: US/10/437,963

NUMBER OF SEQ ID NOS: 2003-05-14

SEQ ID NO 23894

LENGTH: 1693

TYPE: Num
                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Clone ID: PAT_MRT4530_2892C.1 US-10-437-963-23894
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US-10-437-963-23894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Clone ID: MRT4577_77825C.1 US-10-425-115-160357
                                                                                                                                                                                                                                                                                                                                      Query Match 70.7%;
Best Local Similarity 81.9%;
Matches 190; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 23894, Application US/10437963
Publication No. US20040123343A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 213; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               266 AGTCCATCTTCACACACCGGTTGGACAAGGGTTATTGTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14 GTAGAATATGTTAGTGGCTCCTATGATAGGGTGGAAGGATTTGAGTTATTGAATGAGGCA 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            74 ATCTCTGAGTATGAGACTTCAGAAAACAATGACTCGGGAAGCTACCGCAGATTATTTTAT 133
TCATATTGCATGAGTCCATCTTCACACACGGTTGGACAAGGGTTATTGTTG 233
                                                                                                 AGATTATTTTATTTGGCATTGCCTCCATCAGTCTTACCCATCAGTATGCGAGATGATAAGA 181
                                                                                                                                                    TTARATRACGCCATTTCAGAGAATGAAACTTCGAAGAACAACAAACCAGGAAGCTCACGT 1071
                                                                                                                                                                                  TTGAATGAGGCAATCTCTGAGTATGAGACTTCAGAAAACAATGACTCGGGAAGCTACCGC 121
                                                                                                                                                                                                                                          TTTTTGCAATTAATTAAATATGTTAGTGGTTCCTACAATAGCGGGGAAGGATTTGCATCA 1011
                                                                                                                                                                                                                                                                        TTTTTGCAGTTAGTAGAATATGTTAGTGGCTCCTATGATAGGGTTGGAAGGATTTGAGTTA 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TTGGCATTGCCTCCATCAGTCTACCCATCAGTATGCGAGATGATAAGATCATATTGCATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AGTCCATCTTCACACACCGGTTGGACAAGGGTTATTGTTG 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATCTCTGAGTATGAAACTTCAAAAAACAATGACTCGGGAAGCTACCGCAGATTATTTTAT 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GTAAAATATGTTAGTGGCTCCTATGATAGGGTGGAAGGATTTGAGTTATTAAATAAGGCA 145
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                                                                                                                                                                                                                                                                                                                                Score 164.8; DB 18; Length Pred. No. 3.4e-43; 0; Mismatches 42; Indels
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Pred. No. 1.3e-57;
0; Mismatches 7; Indels 0;
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604

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APPLICANT: Zhou, Yihua

APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

FILE REFERENCE: 38-21(53222)B

CURRENT APPLICATION NUMBER: US/10/425,115

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 369326

SEQ ID NO 174626

LENGTH: 749

TYPE: ""
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; Sequence 91761, Application US/10437963
; Publication No. US20040123343A1
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US-10-425-115-174626
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LENGTH: 5580
TYPE: DNA
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APPLICANT:
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OTHER INFORMATION: Clone ID: MRT4577_9083C.1
-10-425-115-174626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
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TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
OTHER INFORMATION: Clone ID: PAT_MRT4530_90304C.1
                                    TYPE: DNA
ORGANISM: Zea mays
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4117 TTTTTGCAATTAATTAATATGTTAGTGGTTCCTACAATAGCGGGAAGGATTTGCATCA 4176
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Wu, Wei
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Barbazuk, Brad
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Pred. No. 5.9e-43;
0; Mismatches 42;
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                                                     RESULT 8
US-10-45-115-48185
; Sequence 48185, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
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CURRENT FILING DATE: 2004-01-29
RUMBER OF SEQ ID NOS: 63128
SEQ ID NO 12821
LENGTH: 713
TYPE: DNA
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Publication No. US20040172684A1
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Best Local Similarity
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Best Local Similarity
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APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OP INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OP INVENTION: Plants and Uses Thereof For Plant Improvement
FILE REFERENCE: 38-21(53535)B
                    APPLICANT: La Rosa, Thomas J. APPLICANT: Kovalic, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Sorghum bicolor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
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    Zhou, Yihua
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78.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 138; DB 18;
Pred. No. 1.4e-34;
0; Mismatches 45;
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Pred. No. 1.5e-35;
0; Mismatches 40;
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503

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US-10-739-930-5153

Sequence 5153, Application US/10739930

; Bedlication No. US20040216190A1

; GENERAL INFORMATION:

; APPLICANT: KOVALIC, David K.

; TITLE OF INVENTION: NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH

; TITLE OF INVENTION: PLANTS AND USES THEREOF FOR PLANT IMPROVEMENT

; FILE REFERENCE: 38-21(53377) B

; CURRENT APPLICATION NUMBER: US/10/739,930

; CURRENT FILING DATE: 2003-12-18

; NUMBER OF SEQ ID NOS: 11088
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Best Local S
Matches 181
                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                 Query Match
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CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 48185
LENGTH: 1929
TYPE: DNA
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TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Clone ID: TRIAE-23APR03-CLUSTER606_35
                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Triticum aestivum FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                         2 ID NO 5153
LENGTH: 1924
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Clone ID: MRT4577_143951C.1
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                                                                                                        515 CTGAACAAGGAATATCAGATTATGAGATGTCA----AACAACTCAGGAAGCTCCCGT
                                                                                                                                                                                         455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   671
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                                                                                                                                                       62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      497 TTTTTGCAACTAATAAAATATGTCAGTGGTTCCTATGATACTGGAGAAGGATTTGAGAAA 556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      181;
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Similarity 76.3%;
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TCATATTGCATGAGTCCATCTTCACACACCGGTTGGACAAGGGTTATTGTTG 233
                                                                  AGATTATTTTATTTGGCATTGCCTCCATCAGTCTACCCATCAGTATGCGAGATGATAAGA 181
                                                                                                                                                                                                          TTTTTGCAGTTAGTAGAATATGTTAGTGGCTCCTATGATAGGGTTGGAAGGATTTGAGTTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AGATTATTTATTTGGCATTGCCTCCATCAGTCTACCCATCAGTATGCGAGATGATAAGA 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGGCTCTTTATTTGGCATTGCCTCCATCTGTCTACCCTTCAGTGTGCAAAATGATCAGA
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78.0%;
                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                       Score 131.6; DB 1
Pred. No. 2.8e-32;
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Pred. No. 2.2e-34;
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                                                                                                                                                                                                                                                        49; Indels
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; LENGTH: 1900
TYPE: DNA
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US-10-437-963-509
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                                                                                                                                                                                                      Sequence 508, Application US/10437963 Publication No. US20040123343A1 GENERAL INFORMATION:
                                                                                                    APPLICANT:
APPLICANT:
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APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
                                                                                                                                                  APPLICANT: La Rosa, Thomas J. APPLICANT: Kovalic, David K. APPLICANT: Zhou, Yihua
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                                                                                      APPLICANT:
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NAME/KEY: unsure
LOCATION: (1)..(1900)
OTHER INFORMATION: unsure at all n locations
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                                                                              Wu, Wei
Boukharov, Andrey
Barbazuk, Brad
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Pred. No. 2.5e-26;
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US-10-425-114-9043
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; OTHER INFORMATION: Clone ID: PAT_MRT4530_100463C.1
US-10-437-963-508
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Publication No. US20040034888A1
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NUMBER OF SEQ ID NOS:
SEQ ID NO 508
LENGTH: 2583
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APPLICANT:
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LENGTH: 1844
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APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
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ORGANISM: Oryza sativa
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  ATCATATTGCATGAGTCCATCTTCACACACCGGTTGGACAAGGGTTATTGTTG: 233
                                                                              CAGATTATTTTATTTGGCATTGCCTCCATCAGTCTACCCATCAGTATGCGAGATGATAAG 180
                                                                                                                      GTTGGATAAAGAGATTTCAGAGCATGAATATTTGAAAAACAGTGTTGAGGGTTTATCTCG
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                                         GAGGCTTTTCTATCTTGCTCTTCCACCTTCAGTATATCCATCAGTTTGCAAGATGATCAA
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Kovalic, David K.
Screen, Steven E
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75.1%;
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Pred. No. 2.9e-26;
0; Mismatches 46
                                                                                                                                                                                                                                                                                              Score 100.2; DB 1
Pred. No. 5.3e-22;
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RESULT 14
US-10-424-599-81077
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US-10-425-114-29479
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; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_44228C.1
US-10-424-599-81077
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CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 29479
LENGTH: 1854
TYPE: DNA
ORGANISM: Glycine max
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                                                                                                     SEQ ID NO 81077
LENGTH: 1947
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Best Local Similarity
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APPLICANT:
                                                                                                                                                                                                              APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
                                                                                                                                                                                                                                                                                                                        APPLICANT: La Rosa Thomas J APPLICANT: Kovalic David K
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TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
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                                                                                                                                                 NUMBER OF SEQ ID NOS: 285684
                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
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                                                            ORGANISM: Glycine max
                                                                                       TYPE: DNA
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Screen, Steven E
Tabaska, Jack E
                                                                                                                                                                                                                                                                                                   Zhou Yihua
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Pred. No. 5.3e-22;
0; Mismatches 83
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RESULT 15
US-10-425-115-142724
; Sequence 142724, Application US/10425115
; Publication No. US20040214272A1
; GENERAL IMFORMATION.
; APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Chou, Yihua
; APPLICANT: Chou, Yihua
; APPLICANT: Coo, Yongwei
TITLE OF INVENTION. Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT APPLICATION UNMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 142724
; LENGTH: 612
TYPE: DNA
; ORGANISM: Zea mays
; OTHER TWENTION:
Search completed: May 4, 2005, 21:11:26 Job time: 266.351 secs
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                                                                                                                                                                                                                                                                                                                                                                                 Query Match 39.3%; Score 91.6; DB 18; Length Best Local Similarity 74.4%; Pred. No. 2.1e-19; Matches 131; Conservative 0; Mismatches 39; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 43.0%; Score 100.2; DB 17; Length 1947; Best Local Similarity 64.4%; Pred. No. 5.4e-22; Matches 150; Conservative 0; Mismatches 83; Indels 0;
                                                                                                 554
                                                                                                                                                                                   500 CTGAACAAGGAAATATCAGATTATGAGATGTC----CAACAACTCGGGAATCTCCCGC 553
                                                                                                                                                                                                                                                                                  440 TTCTTGCAATTGATAAAATATGTCAGTGGTTCCTATGACAGCGGAGAAAGTTTTGAAAAA 499
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                                                                                                                                                                                                                          62 TTGAATGAGGCAATCTCTGAGTATGAGACTTCAGAAAACAATGACTCGGGAAGCTACCGC 121
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                                                                                                                           AGATTATTTATTTGGCATTGCCTCCATCAGTCTACCCATCAGTATGCGAGATGAT 177
                                                                                    AGGCTATTTTATTTGGCATTGCCTCCATCTGTTTACCCTTCACTGGGCAAAATGAT 609
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GenCore version 5.1.6
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Title:
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Listing first 45 summaries
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

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5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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US-09-710-279-3602
US-09-107-532A-1949
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53 AGGGATGGAATCGCATAATATTTGAGAAGCCATTTGGCTTTGATGCACTTTCTTCCCATA 112

Query Match 23.3%; Score 61.4; DB 4; Length Best Local Similarity 52.1%; Pred. No. 1.3e-08; Matches 110; Conservative 0; Mismatches 101; Indels	US-09-557-884-1/c US-09-557-884-1/c Sequence 1, Application US/09557884 Patent No. 6506581 GENERAL INFORMATION: APPLICANT: Fleischmann et al. TITLE OF INVENTION: The Nuclectide sequence of the Haemophilus influenzae Rd TITLE OF INVENTION: Thereof, and Uses Thereof NUMBER OF SEQUENCES: 1 CORRESPONDENCE ADDRESS: 1 CORRESPONDENCE ADDRESS: 1 COUNTRY: USA ZIP: 20850 COUNTRY: USA ZIP: 20850 COMPUTER READABLE FORM: MEDIUM TYPE: 3 1/2 inch diskette COUNTRY: USA ZIP: 20850 COMPUTER READABLE FORM: MEDIUM TYPE: 3 1/2 inch diskette COMPUTER READABLE FORM: MEDIUM TYPE: 3 1/2 inch diskette COMPUTER READABLE FORM: MEDIUM TYPE: 3 1/2 inch diskette COUNTRY: USA ZIP: 20850 COMPUTER READABLE FORM: MEDIUM TYPE: 3 1/2 inch diskette COUNTRY: USA ZIP: 20850 COMPUTER READABLE FORM: MEDIUM TYPE: 3 1/2 inch diskette COUNTRY: USA ZIP: 20850 COMPUTER READABLE FORM: MEDIUM TYPE: 3 1/2 inch diskette COUNTRY: USA ZIP: 20850 COMPUTER READABLE FORM: MEDIUM TYPE: 3 1/2 inch diskette COUNTRY: USA ZIP: 20850 COMPUTER: ASCII Text CURRENT APPLICATION UNMER: US/99/557,884 FILING DATE: 25-Apr-2000 CLASSIFICATION: UNMER: US/99/557,884 FILING DATE: 25-Apr-2000 CLASSIFICATION UNMER: US/99/557,884 FILING DATE: 25-Apr-2000 CLASSIFICATION UNMER: US/99/557,884 FILING DATE: 25-Apr-2000 FILING DATE: 25-Apr-2000 CLASSIFICATION UNMER: US/99/557,884 FILING DATE: 301-309-850,476,102 FILING DATE: JUN-5-1995 ATTORNATION FOR SEO ID NO: 1: SEQUENCE CHARACTERISTICS: LENGTH: 1830121 base pairs TELEPHONE: 301-309-850,4 TELEPH	C 28 42 16.0 3376 4 US-09-710-279-4055 29 39.6 15.1 6995 4 US-09-531-266-1 30 37.4 14.2 1461 3 US-08-445-463B-5 31 37.4 14.2 1461 3 US-08-044-8570-5 32 37.4 14.2 1461 3 US-08-044-8570-7 35 37.2 14.1 1455 5 PCT-US94-03437-7 35 37.2 14.1 1457 5 PCT-US94-03437-7 37.2 14.1 1457 5 PCT-US94-03437-7 37.2 14.1 1571 3 US-08-445-463B-3 37.2 14.1 1571 3 US-08-445-464C-3 37.2 14.1 1571 3 US-08-044-8570-3 39 37.2 14.1 1696 1 US-07-737-071A-1 40 37.2 14.1 1696 1 US-08-022-096-1 41 37.2 14.1 1696 3 US-08-445-463C-7 42 37.2 14.1 1696 3 US-08-445-463C-7 43 37.2 14.1 1696 3 US-08-445-463C-7 44 36.4 13.8 34266 4 US-09-949-016-13250 45 36.2 13.8 2853 4 US-09-620-312D-463
.h 1830121; .ls	Genome, Fragments	Sequence 1, Appli Sequence 5, Appli Sequence 5, Appli Sequence 5, Appli Sequence 5, Appli Sequence 7, Appli Sequence 3, Appli Sequence 3, Appli Sequence 1, Appli Sequence 1, Appli Sequence 7, Appli Sequence 13250, A Sequence 13250, A

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US-09-643-990A-1
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                                                              Best Local Similarity 52. Matches 110; Conservative
                                                                                                  Query Match
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GENERAL INFORMATION:
                                                                                                                                                                                                                                             TELEFAX: 310-309-8439 INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: 3 1/2 inch diskette
COMPUTER: Dell Pentium
OPERATING SYSTEM: MS DOS v6.22
SOPTWARE: ASCII Text
SOPTWARE: ASCII Text
APPLICATION NUMBER: US/09/643,990A
FILING DATE: 23-Aug-2000
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/487,429
FILING DATE: 1995-06-07
APPLICATION NUMBER: 08/426,787
FILING DATE: 1995-04-21

PRICE DATE: 1995-04-21
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                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Kenley K. Hoover
REGISTRATION NUMBER: 40,302
REFERENCE/DOCKET NUMBER: PB1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 301-610-5790
                                                                                                                                                  SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           J. Craig Venter
J. Craig Venter
TITLE OF INVENTION: The Nucleotide sequence of the Haemophilus influenzae Thereof, and Uses Thereof
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ADDRESSEE: Human Genome Sciences,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
53 AGGGATGGAATCGCATAATATTTGAGAAAGCCATTTGGCTTTGATGCACTTTCTTCCCATA 112
                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                        LENGTH: 1830121 base pairs
                                                                                                                                                                       COPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 9410 Key West Avenue CITY: Rockville, STATE: MD
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Owen White
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                                                                          23.3%;
                                                            0;
                                                       Score 61.4; DB 4;
Pred. No. 1.3e-08;
0; Mismatches 101;
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                                                                                           Length 1830121;
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RESULT 4
US-09-134-000C-2369
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TITLE OF INVENTION: Anti-Bacterial Vaccine Compositions
FILE REFERENCE: 28341/00435
CURRENT APPLICATION NUMBER: US/09/809,665A
CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: 60/153,453
PRIOR FILING DATE: 1999-09-10
PRIOR APPLICATION NUMBER: 60/128,689
PRIOR APPLICATION NUMBER: 60/128,689
PRIOR FILING DATE: 1999-04-09
PRIOR FILING DATE: 1999-04-09
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
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GENERAL INFORMATION:
APPLICANT: LYMN DOUGETTE-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: ENTEROCOCCUS FAECALIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 032796-032
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SEQ ID NO 7
LENGTH: 6132
TYPE: DNA
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Best Local Similarity
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Patent No. 6790950
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LOCATION: (4032)..(4727)
OTHER INFORMATION: devB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 340:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: KIMMERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR PPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No. 6703492
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 6812
SOFTWARE: PatentIn version 3.1
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TYPE: DNA
ORGANISM: Enterococcus faecalis
                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: nucleic acid sequence
                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
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225
                                                                        165
                                                                                                                                                                                                                                                             114; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           53 AGGGATGGAATCGCATAATATTTGAGAAGCCATTTGGCTTTGATGCACTTTCTTCCCATA 112
                                                                                                                                                                                                                        45 GACCCAGAAGGGATGGAATCGCATAATATTTGAGAAGCCATTTGGCTTTGATGCACTTTC 104
                                                                                                                                                                                                                                                                               Similarity
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AGNTTTTGAGCCACCTTNGAGNNGNACNT 253
                                   TGATCACTATTTAGGTAAAGAAATGATTCAAAATATTGAACGTCTACGATTTTGAAAACAC 1492
                                                                                                           T--GCAAAAATTAAATAATCAAATAAGAAAGTCTTTTAAGAAGAAAAAATATTTGGAAT
                                                                                                                                                TTCCCATAGGCTGACACTATCTTCTTTCAAACTTTCAGGAAAAGCAAATATATAGAAT 164
                                                                      TGANCATCTACTAGGAAGGAATCNCAGTNAAAATCCTNCAGGTTTAAGGGTTTCAAANNN 224
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                                                                                                                                                                                                                                                                             Score 54; DB 4;
Pred. No. 1.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 58.2; DB 4;
Pred. No. 5.3e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 4;
                                                                                                                                                                                                                                                       93;
                                                                                                                                                                                                                                                                                              Length 3015;
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                                                                                                                                                                                                                                                                                                                                                                        synthetic
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                                                                                                                                                                                                                                                           Gaps
                                                                                                           1432
                                                                                                                                                                                   1374
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                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-107-532A-1949
                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-710-279-3602/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-710-279-3602
                                                                                                                                                                                                                                                                                                                                                                    Patent No. 6583275
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. SEQ ID NO 3602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 6703492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: PU3480US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2228
                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1493 TATATTTGAACCACTTTGGAATAATAAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        165
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                                                                                                                                                               COUNTRY: USA
ZIP: 02354
                                                                                                                                                                                                                          CITY: Waltham
                                                                                                                                                                                                                                          STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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Sequence 1949, Application US/09107532A
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PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: KIMMERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/107,532A
FILING DATE: 30-Jun-1998
                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
COMPUTER: PC
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2168 TATATTTGAACCACTTTGGAATAATAAGT 2140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2286 T--GCAAAAATTAAATAATCAAATAAGAAGTCTTTTAAGAAGAAAAAAATATTTGGAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2346 GAACGTAAAAGGGGGTGATGCGAATAATTATAGAAAAACCATTTGGTGATGATTTAAAATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          45 GACCCAGAAGGGATGGAATCGCATAATATTTGAGAAAGCCATTTTGGCTTTTGATGCACTTTC
                                                                                           COMPUTER: PC
OPERATING SYSTEM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TGANCATCTACTAGGAAGGAATCNCAGTNAAAATCCTNCAGGTTTAAGGGTTTCAAANNN 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGNITITGAGCCACCTINGAGNNGNACNI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TICCCATAGGCTGACACAATATCTTCTTTCAAACTTTCAGGAAAAGCAAATATATAGAAT 164
                                                                                                                                                                                                                                                                           STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: GENOME THERAPEUTICS CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TGATCACTATTTAGGTAAAGAAATGATTCAAAATATTGAACGTCTACGATTTTGAAACAC 2169
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                                                                                                                                                                                                                                                                                                                              100 Beaver Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Description of Artificial Sequence: synthetic nucleic acid sequence
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Pred. No. 1.8e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            253
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RESULT 8
US-08-956-171E-53
; Sequence 53, Application US/08956171E
; Patent No. 6593114
; GENERAL INFORMATION:
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; LOCATION: (B) LOCATION 1...1548
; SEQUENCE DESCRIPTION: SEQ ID NO: 1949:
US-09-107-532A-1949
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (781)893-8277
INFORMATION FOR SEQ ID NO: 1949:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 60/085,598
FILING DATE: 14 May 1998
APPLICATION NUMBER: 60/051571
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Arinitallo, Pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-012
TELEPHONE: (781)893-5007
TELEPHONE: (781)893-5007
ZIP: 20850
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
                                                                                                                                                                                                                   Gil H. Choi
Patrick S. Dillon
Craig A. Rosen
Steven C. Barash
Michael R. Fannon
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
NUMBER OF SEQUENCES: 5256
                                                                                                                                                    CORRESPONDENCE ADDRESS:

ADDRESSEE: Human Genome Sciences,
STREET: 9410 Key West Avenue
CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: circular MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     599
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          479 CTGATTACCAGTCTGCTTTTGAATTGAACCAGCAAATCAGAGAAGTATTCCCCAGAACAAG 538
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92 TIGATGCACTTICTTCCCATAGGCTGACAÇAATATCTTCTTCAAACTTTCAGGAAAAGC 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     32 CAAGCAGTGCTCAGACCCAGAAGGGATGGAATCGCATAATATTTTGAGAAGCCATTTGGCT 91
                                                                                                                STATE: Maryland COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GTTTTGCCAATAATATTTTTGAGGCTCAATGGAACAATCGCTATATCGACAA 650
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGGTTTCAAANNNAGNTTTTGAGCCACCTTNGAGNNGNACNTNNNNNNGANNA 263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATATTTTCAGGATTGATCATTATTTAGGAAAAGAAATGATCCAAAACATCTCAGCTATCC 598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CACAAAATATTATGACAAAAAATGGCTTCAATCGTTTGATCATAGAAAAACCGTTCGGTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Enterococcus faecium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 1548 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                          Charles Kunsch
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48.3%; Pred. No. 5.9e-07;
tive 0; Mismatches 120; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 1548;
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US-08-781-986A-53
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APPLICATION NUMBER: 60/009,861
FILING DATE: January 5, 1996
APPLICATION NUMBER: 08/781,986
FILING DATE: January 3, 1997
ATTORNEY/AGENT INFORMATION:

NAMB: MARK J. Hyman

REGISTRATION NUMBER: 46,789
REFERENCE/DOCKET NUMBER: PB248P1
TELECOMMUNICATION INFORMATION:
TELEPAN: (301) 309-314-1224
TELEPAX: (301) 309-8439
INFORMATION FOR SEQ ID NO: 53:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 53, Application US/08781986A Patent No. 6737248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 103;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Charle
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                    MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/781,986A
                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Charles Kunsch
TITLE OF INVENTION: Staphylococcus aureus
NUMBER OF SEQUENCES: 5255
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                     STREET: 9410 Key West Avenue
CITY: Rockville
STATE: Maryland
COUNTRY: USA
                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: CURRENT APPLICATION DATA:
    APPLICATION UNMBER: US/08/956,171E
    FILING DATE: 20-Oct-1997
    CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          934 TATTTÄGGÄÄAAGÄCATGGTTCÄÄÄÄTATCGAGGTATTÄCGTTTTGCGÄÄTGCGATGTTT 993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               874 GCATTAÀACAÀTCÀAATTCGTAAÀTCATTTAÀAGÀAGAAGAÀATTTATCGTATTGACCAC 933
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         112 AGGCTGACACAATATCTTCTTTCAAACTTTCAGGAAAAGCAAATATATAGAATTGANCAT 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      814 AAAGGATTTAAACGCCTTGTTATCGAAAAACCATTCGGTAGTGATTTAAAATCAGCCGAA 873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             994 GAACCATTATGGAATAACAAAT 1015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       232 GAGCCACCTTNGAGNNGNACNT 253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              52 AAGGGATGGAATCGCATAATATTTGAGAAGCCATTTGGCTTTGATGCACTTTCTTCCCAT 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTACTAGGAAGGAATCNCAGTNAAAATCCTNCAGGTTTAAGGGTTTCAAANNNAGNTTTT 231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 16592 base pairs TYPE: nucleic acid STRANDEDNESS: double TOPOLOGY: linear
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ilarity 51.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                          Human Genome Sciences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 51.4; DB 4;
Pred. No. 2.8e-06;
O; Mismatches 99;
                                                                                                                                                                                                                                                                                                                                                                                                                  Polynucleotides and Sequences
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RESULT 10
US-08-714-918-3/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-781-986A-53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 6037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Sir
Matches 103;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (301) 309-8512 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTONNELL...

NAME: Benson, Bob
REGISTRATION NUMBER: 30,446
RECISTRATION NUMBER: PB24/
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                             CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Sun, Dongxu
TITLE OF INVENTION: STAPHYLOCOCCI
TITLE OF INVENTION: TARGET GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                            CURRENT APPLICATION DATA: US
                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                        COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. DOS 5.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                               FILING DATE:
                                                                                                                           SOFTWARE:
                 APPLICATION NUMBER: 60/009,102 FILING DATE: December 22, 1995
                                                                                                                                                                                                                                      COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CATION NUMBER:
                                                                                                                                                                                                                     90071-2066
                                                                                                                                                                                                                                                 Los Angeles
California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/08714918
                                                                                                                                                                                                                                                                                                                                             SEQUENCES:
                                                                                                                                                                                                                                                                                Suite 4700
                                                                                                                                                                                                                                                                                                  633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                        Malouin, Francois
Martin, Patrick K.
Schmid, Molly B.
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                                                                                                                           Word Perfect 5.1
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                                                                            September 13,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                double
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                                                                                          US/08/714,918
60/003,798
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Pred. No. 2.8e-06;
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                                                                               1996
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US-09-265-315-3/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: March 9,
                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
APPLICANT:
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TYPE: n
                                                                                          OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
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   APPLICATION NUMBER:
                                  CLASSIFICATION
                                                                                                                         COMPUTER:
                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE:
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                                                                                                                                                                                        90071-2066
                                                                                                                         IBM Compatible
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sequence 3, Application US/09265315 Patent No. 6187541
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                TITLE OF INVENTION: ITTILE OF INVENTION: ITTILE OF INVENTION: ITTILE OF SEQUENCES:
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TELECOMMUNICATION INFORMATION:
                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 Weet Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: September : ATTORNEY/AGENT INFORMATION
                                                                          STATE: Calif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1245 GAACCATTATGGAATAACAAAT 1224
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1365 GCATTAAACAATCAAATTCGTAAATCATTTAAAGAAGAAGAAATTTAYCGTATKGACCAC 1306
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Martin, Patrick K.
Schmid, Molly B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                     Benton, Bret
Lee, Ving J.
Malouin, Francois
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                                                                                                                                                                                                                                                                                                      Dongxu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19.2%;
50.0%;
                                                                                                                                                                                                                                                           METHODS OF SCREENING FOR COMPOUNDS ACTIVE ON STAPHYLOCOCCUS AUREUS
                                                                                                                                                                                                                                    TARGET GENES
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Pred. No. 2.1e-06;
1.44 Mb
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US/09/265,315

DOS 5.0

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US-09-265-315-3/c
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SEQUENCE CHARACTERISTICS:
LENGTH: 2494 hard
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Best Local
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FILING DATE: September 15, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 240/247
TELECOMMUNICATION INFORMATION:
TELEPHONE: / 1211
                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
COMPUTER: IBM COmpatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: METHODS OF SCREENING FOR COMPOUTITIES OF INVENTION: ACTIVE ON STAPHYLOCOCCUS AUREUS TITLE OF INVENTION: TARGET GENES
                                                                                                                                                                                                                                                                                                                                                 STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: LOS Anno?
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                 COMPUTER: IBM CON
OPERATING SYSTEM:
          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                          CITY: Los Angeles
STATE: California
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APPLICATION NUMBER:
FILING DATE: Decembe
APPLICATION NUMBER:
FILING DATE: Septemb
                                                                                                                                                                                                                                                                                               COUNTRY:
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STRANDEDNESS: sing
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TELEFAX: (213) 955-0440
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                                                                                                                                                                                                                                                                    90071-2066
                                     TING SYSTEM: IBM P.C. ARE: Word Perfect 5.1 APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GCATTAAACAATCAAATTCGTAAATCATTTAAAGAAGAAGAAATTTAYCGTATKGACCAC 1306
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Lee, Ving J.
Malouin, François
Martin, Patrick K.
Schmid, Molly B.
                                                                                                                                                                                                                                                                                               U.S.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sun, Dongxu
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UMBER: 60/009,102
December 22, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      single
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US/09/265,315
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Pred. No. 2.1e-06;
2; Mismatches 99;
                                                                                                 DOS 5.0
                                                                                                                                                                                             1.44 Mb
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GENERAL INFORMATION:
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Best Local S
Matches 101
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: March 9, 1999
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/714,918
FILING DATE: September 13, 1996
APPLICATION NUMBER: 60/003,798
PILING DATE: December 22, 1995
APPLICATION NUMBER: 60/003,798
         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                    APPLICANT: Lee, Ving J.
APPLICANT: Malouin, François
APPLICANT: Martin, Fatrick K.
APPLICANT: Schmid, Molly B.
APPLICANT: Schmid, Molly B.
APPLICANT: Sun, Dongxu
TITLE OF INVENTION: METHODS OF SCREENING FOR COMPOUNDS
TITLE OF INVENTION: ACTIVE ON STAPHYLOCOCCUS AUREUS
NUMBER OF SEQUENCES: 111
                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 240/247
TELECOMMUNICATION INFORMATION:
                                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: September 15, 1995
ATTORNEY/AGENT INFORMATION:
                                                                   COUNTRY: U.S.A. ZIP: 90071-2066
COMPUTER:
                                                                                                                                                                              ADDRESSEE:
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nucleic acid
DEDNESS: single
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IBM
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Pred. No. 2.1e-06;
2; Mismatches 99;
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RESULT 14
US-09-528-709-3/c
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                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3, Application US/09528709 Patent No. 6630303 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 101;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 19.2%;
Best Local Similarity 50.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 2494 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: December 22, 1995
APPLICATION NUMBER: 60/003,798
FILING DATE: September 15, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Word Perfect CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 2494 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 240/248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 60/009.
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FILING DATE: March 9, 1999
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                                                                                                                       NUMBER OF SEQUENCES: 111
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
COMPUTER
                                                                                                                                                                                                               TITLE OF INVENTION: STAPHYLOCOCCUS AUREUS ANTIBACTERIAL TARGET GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1425 AAAGGATTTAAACGCCTTGTTATCGAAAAACCATTCGGTAGTGATTTAAAATCAGCCGAA 1366
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAACCATTATGGAATAACAAAT 1224
                                                          CITY: Los Angeles
STATE: California
                  COUNTRY: U.S.A. ZIP: 90071-2066
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Lee, Ving
Malouin, Francois
Martin, Patrick K.
Schmid, Molly B.
                                                                                                     Suite 4700
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Perfect 5.1
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mber 13, 1996
60/009,102
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Pred. No. 2.1e-06;
2; Mismatches 99;
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US-09-527-745-3/c
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                                                                                                                                                                                                                                                                         Sequence 3, Application Use Patent No. 6638718
GENERAL INFORMATION:
APPLICANT: Benton,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 101; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 2494 base pairs
TYPE: nucleic acid
STRANDENESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/714,918
FILING DATE: September 13, 1996
APPLICATION NUMBER: 60/009,102
FILING DATE: December 22, 1995
APPLICATION NUMBER: 60/003,798
FILING DATE: September 15, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1305
                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                     NUMBER OF SEQUENCES: 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                     1245 GAACCATTATGGAATAACAAAT 1224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1365 GCATTAAACAATCAAATTCGTAAATCATTTAAAGAAGAAGAAATTTAYCGTATKGACCAC 1306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1425 AAAGGATTTAAACGCCTTGTTATCGAAAAACCATTCGGTAGTGATTTAAAATCAGCCGAA 1366
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   172 CTACTAGGAAGGAATCNCAGTNAAAATCCTNCAGGTTTAAGGGTTTCAAANNNAGNTTTT 231
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TATTTAGGAAAAGACATGGTTCAAAATATCGAGGTATTACGTTTTTGCGAATGCGATGTTT 1246
                                                                                                                                                                                                                                                                                                                                        Application US/09527745
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                         INVENTION: STAPHYLOCOCCUS AUREUS ANTIBACTERIAL TARGET GENES
                                                                                                                                                                                  Lee, Ving
Malouin, Francois
Martin, Patrick K.
Schmid, Molly B.
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CLASSIFICATION: 4.4

PRIOR APPLICATION NUMBER: US/08/714,918
FILING DATE: September 13, 1996
APPLICATION NUMBER: 60/009,102
FILING DATE: December 22, 1995
APPLICATION NUMBER: 60/003,798
FILING DATE: September 15, 1995
APPLICATION NUMBER: 60/003,798
FILING DATE: September 15, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 222/005
TELECOMMUNICATION: NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 222/005
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEYA: 67-3510
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 2494 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 3:
US-09-527-745-3
Search completed: May 4, 2005, 12:27:36 Job time : 81.9588 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 19.2%; Score 50.6; DB 4; Length 2494; Best Local Similarity 50.0%; Pred. No. 2.1e-06; Matches 101; Conservative 2; Mismatches 99; Indels 0;
                                                                                                                                                                         1305 TATTTAGGAAAAGACATGGTTCAAAATATCGAGGTATTACGTTTTGGGAATGCGATGTTT 1246
                                                                                                                                                                                                                                                                1365 GCATTAÀACAATCAAATTĊGTAAATCATTTAÀAGAAGAAGAAATTTAYCGTATKGACCAC 1306
                                                                                                                                                                                                                                                                                                                                                          1425 AAAGGATTTAAACGCCTTGTTATCGAAAAACCATTCGGTAGTGATTTAAAATCAGCCGAA 1366
                                                                                         1245 GAACCATTATGGAATAACAAAT 1224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                232 GAGCCACCTTNGAGNNGNACNT 253
                                                                                                                                                                                                                                                                                                     CTACTAGGAAGGAATCNCAGTNAAAATCCTNCAGGTTTAAGGGTTTCAAANNNAGNTTTT 231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/527,745
FILING DATE: 17-Mar-2000
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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Minimum DB seq
Maximum DB seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Run on:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           length: 0
length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  May 4, 2005, 12:06:42; Search time 298.387 Seconds (without alignments) 5365.716 Million cell updates/sec
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Gapop 10.0 , Gapext 1.0
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263
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                                                                                                                                                                                                                                                                                                                                                                                            gaagcacttttggatgttgc.....agnngnacntnnnnnganna 263
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Copyright (c) 1993 - 2005 Compugen Ltd.
/ Ggn2_6/ptodata/1/pubpna/US09C_PUBCOMB.seq:*
/ Cgn2_6/ptodata/1/pubpna/US09. NEW PUB.seq:*
/ Cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
/ Cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
/ Cgn2_6/ptodata/1/pubpna/US10C_PUBCOMB.seq:*
/ Cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*
                                                                                                                                                                                                                                                                                                               Cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
Cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
Cgn2_6/ptodata/1/pubpna/US09A_PUBCOMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

, ,	Result
10087654321	· ir
235 215.2 215.2 215.3 175.8 123.2 71.8 64.2 61.8 61.4 61.4	Score
223.554.4 23.554.3 23.554.3 3.554.3	Query Match
1680 2218 237 260 2218 1577 1482 1501 1485 1485	Query Match Length DB
17 17 10 10 17 17 17	DB
US-10-425-114-8171 US-10-424-599-48573 US-09-237-183A-2682 US-09-237-183A-2684 US-10-424-599-3107 US-10-282-122A-9150 US-10-282-122A-9150 US-10-282-122A-9150 US-09-815-242-6958 US-10-282-122A-22036	ID
Sequence 8171, Ap Sequence 48573, A Sequence 2682, Ap Sequence 2684, Ap Sequence 3107, Ap Sequence 3107, Ap Sequence 9150, Ap Sequence 9150, Ap Sequence 6558, Ap Sequence 6280, Ap	Description

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1530	1767	1534	1482	1320	972	3011208	319630	2319	2812	2328	1396	1491	840	1800	1800	32768	1524	1521	1476	1731	1731	1464	6132	6132	1494	2632	2274	2274	2249	2232	1782	1782	1830121	
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US-10-369-493-26367	+0.000 +000 0+00	116-10-282-1228-21509	US-10-369-493-25835	US-10-369-493-25834	US-10-282-122A-35419	US-10-398-221-2058		US-10-739-930-5533	24-599-3	US-10-425-114-6121	US-10-424-599-31276	US-10-282-122A-31151	US-10-425-115-109727	US-09-938-842A-1621	US-09-938-842A-1621	US-09-070-927A-128	US-09-815-242-6547	-282-122A-2109	US-10-282-122A-24529	US-09-938-842A-156	US-09-938-842A-156	US-10-369-493-40267	US-10-854-299-7	US-09-809-665A-7	US-10-282-122A-12546	US-10-425-115-162957	US-10-425-114-830	114-	US-10-425-114-20264	US-10-425-114-16607	US-09-938-842A-2455	US-09-938-842A-2455	US-10-158-865-1	
Sequence 26367, A	CACCE CONTRACT	21500	25835,	Sequence 25834, A	,	Sequence 2058, Ap	/, Appı	Sequence 5533, Ap	31		31276,	31151,	10972	e 1621	1621, /	128,	6547		Sequence 24529, A	Sequence 156, App		Sequence 40267, A	7, Appl		12546,	1629		777,		16607,	2455,	455,	-	

ALIGNMENTS

US-10-425-114-8171 ; Sequence 8171, Application US/10425114 ; Publication No. US20040034888A1

GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua

```
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
ITILE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION UNUMER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
INUMBER OF SEQ ID NOS: 73128
SEQ ID NO 8171
LENGTH: 1680
TYPE: DNA
ORGANISM: Glycine max
FEATURE:
OTHER INFORMATION: Clone ID: 700737284_FLI
US-10-425-114-8171

Query Match
Best Local Similarity 90.5%; Pred. No. 1.3e-59;
Matches 238; Conservative 0; Mismatches 25; Indels 0; Gaps 0;
Matches 238; Conservative 0; Mismatches 25; Indels 0; Gaps 0;

1 GAAGCACTTTTGGATGTTGCGTCATGTCTTGCAAGCCCAGAAGGGATGG 501
Db 442 GAAGCACTTTTGGATGTTTGCGTCATGTCTTGCAAGCCCAGAACGGATGG 501

Qy
61 AATCGCATAATATTTGGAAGCCATTTTGGATGTCTTGCATGTCTTCCTAGACCCAGAAGGGATGG 501
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US-09-237-183A-2682
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-424-599-48573
                                                                                                  Sequence 2682, Application US/09237183A Publication No. US20030135870A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5322)B
CURRENT TAPLICATION NUMBER: US/10/424,599
CURRENT TILING DATE: 2003-04-28
SQQ ID NO 48573
LENGTH: 2218
TYPE: DNA
TYPE: DNA
                  APPLICANT: Cheikh, No. US20030135870Aldine
APPLICANT: Fisher, Dane K.
APPLICANT: Liu, Jingdong
TITLE OF INVENTION: Nucleic Acid Molecules And
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Clone ID: PAT_MRT3847_14868C.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Glycine max FEATURE:
                                                                                                                                                                                                                                                 1134
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       954 AATCGCATAATATTTGAGAAGCCATTTGGCTTTGATGCACTTTCTTCCCCATAGGCTGACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     894 GAAGCACTITTGGATGTTGCGTCATGTCTTGCAAGCAGTGCTCAGACCCCAGAAGGGATGG 953
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                                                                                                                                                                                                                                           TGGAGTCGTACTTATATAGATAA 1156
                                                                                                                                                                                                                                                                                                                                               AGGAATCNCAGTNAAAATCCTNCAGGTTTAAGGGTTTCAAANNNAGNTTTTGAGCCACCT 240
                                                                                                                                                                                                                                                                                                                                                                                                                           CAATATCTTCTTCAAACTTTCAGGAAAAGCAAATATATAGAATTGANCATCTACTAGGA 180
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                                                                                                                                                                                                                                                                                TNGAGNNGNACNTNNNNNGANNA 263
                                                                                                                                                                                                                                                                                                                                                                                              CAATATCTTCTTTCAAACTTTCAGGAAAAGCAAATATATAGAATTGATCATCTACTAGGA
                                                                                                                                                                                                                                                                                                                    AGGAATCTCATTGAAAATCTTACAGTTTTAAGGTTTTCAAATCTAGTTTTTGAGCCACTT
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Pred. No. 1.5e-59;
0; Mismatches 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25;
              Other Molecules Associated With
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 2218;
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APPLICANT: Cheikh, No. US20030135870Aldine
APPLICANT: Liu, Jingdong
ITITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With The
TITLE OF INVENTION: Sucrose Pathway
FILE REFERENCE: 38-21(1508)B
CCURRENT APPLICATION NUMBER: US/09/237,183A
CCURRENT FILING DATE: 1999-01-26
PRIOR FILING DATE: 1999-11-24
PRIOR APPLICATION NUMBER: US 60/067,000
PRIOR APPLICATION DATE: 1997-11-24
SEQ ID NO 2684
LENGTH: 260
TYPE: DNA
ORGANISM: Glycine max
US-09-237-183A-2684
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US-09-237-183A-2684
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NAME/KEY: unsure
LOCATION: (1)..(237)
OTHER INFORMATION: unsure at all n locations
US-09-237-183A-2682
                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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Publication No. US20030135870A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches 221;
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Best Local
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LENGTH: 237
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FILE REFERENCE: 38-21(15089)B
CURRENT APPLICATION NUMBER: US/09/237,183A
CURRENT FILING DATE: 1999-01-26
PRIOR APPLICATION NUMBER: US 60/067,000
PRIOR FILING DATE: 1997-11-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 2814
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
177 AGGAAGGAATCNCAGTNAAAATCCTNCAGGTTTAAGGGTTTCAAANNNAGNTTTTGAGCC 236
                                                                                        117
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                                                                                                                                N
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                                       GACACAATATCTTCTTTCAAACTTTCAGGAAAAGCAGATATATAGAATTGATCATCTACT
                                                                   GACACAATATCTTTCTTTCAAACTTTCAGGAAAAGCAAATATATAGAATTGAACATCTACT 176
                                                                                                                                                       ATGGAATCGCATAATATTTGAGAAGCCATTTGGCTTTGATGCACTTTCTTCCCATAGGCT 116
                                                                                                                     ATGGAATCGCATAATATTTGATAAGCCATTTGGCTTTGATGCACTTTCTTCCCCATAGGCT
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                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                          66.8%;
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93.2%;
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; Pred. No. 2.6e-
0; Mismatches
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Pred. No. 4e-54;
0; Mismatches
                                                                                                                                                                                  2.6e-42;
-has 27; Indels
                                                                                                                                                                                                                                             DB 10;
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                                                                                                                                                                                                                                           Length 260;
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US-10-424-599-3107; Sequence 3107, Application US/10424599; Publication No. US20040031072A1
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JS-10-437-963-65487
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LENGTH: 2218
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                                                                                                                                                              GENERAL INFORMATION:
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CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53221)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                               APPLICANT: La Rosa Thomas
APPLICANT: Kovalic David
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
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ORGANISM: Oryza sativa
FEATURE:
      PPLICANT: Cao Yongwei
TTLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TTLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
LE REFERENCE: 38-21 (53223) B
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                                                                                                                                                                                                                                                                                                     TGGAGTCGCACCT 1193
                                                                                                                                                                                                                                                                                                                                          TNGAGNNGNACNT 253
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Barbazuk, Brad
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NUMBER: US/10/424,599
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66.0%;
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                                                                                                                  PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR PPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/230,335
PRIOR APPLICATION NUMBER: 60/230,335
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NUMBER OF SEQ ID NOS:
SEQ ID NO 3107
LENGTH: 1577
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Publication No. US20040029129A1
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                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Identification of Essential Genes in Microorganisms FILE REFERENCE: ELITRA.034A
                                                            PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 2000-09
                                                                                                     PRIOR FILING DATE: 2000-09-06
                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/282,122A
CURRENT FILING DATE: 2003-02-20
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LOCATION: (1)..(157
OTHER INFORMATION:
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ORGANISM: Glycine
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mes 133; Conserv
                                       APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zamudio, Carlos
Malone, Cheryl
Haselbeck, Robert
                                                                                                                                                                                                                                                                                                                                                           Carr, Grant
Yamamoto, Robert
Forsyth, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Wall, Daniel
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Zyskind, Jud
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lio, Carlos
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                                       60/242,578
                                                                               60/230,347
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Pred. No. 8.
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APPLICANT: Oblisen, Kari
APPLICANT: Zyskind, Judith
APPLICANT: Wall, Daniel
APPLICANT: Wall, Daniel
APPLICANT: Trawick, John
APPLICANT: Trawick, John
APPLICANT: Forsyth, R.
APPLICANT: Forsyth, R.
APPLICANT: YAMAMOOTO, Robert
APPLICANT: YOUNG H.
APPLICANT ON WINDER: US/10/282,122A
CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/230,335
PRIOR APPLICATION NUMBER: 60/230,347
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR APPLICATION NUMBER: 60/250,625
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; TYPE: DNA
; ORGANISM: Bacillus anthracis
US-10-282-122A-9984
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PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: 60/267,636
PRIOR FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/269,308
PRIOR FILING DATE: 2001-02-16
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SEQ ID NO 9984
LENGTH: 1482
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               151 CAAATATATAGAATTGANCATCTACTAGGAAGGAATCNCAGTNAAAATCCTNCAGGTTTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAAAGCGGACTTGATAAAACGGATGGAAAACGCCTAATGAATTGAGAAACCGTTCGGG 459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zamudio, Carlo
Malone, Chery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Haselbeck, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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Pred. No. 1.5e-08;
0: Mismatches 97;
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                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 126220
LENGTH: 214
                                                                                                                                                     Matches
                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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Publication No. US20040031072A1
                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: La ROBA Thomas J
APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
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SEQ ID NO 9150
LENGTH: 1501
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                                                                                                                                                                                                                                                                   NAME/KEY: unsure
LOCATION: (1)..(214)
OTHER INFORMATION: unsure at all n locations
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Glycine
                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: 60/267,636
PRIOR FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/269,308
PRIOR FILING DATE: 2001-02-16
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                                223 NNAGNTTTTGAGCCACCTTNGAGNNGNACNTNNNNNGANNA
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        61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        414 GATGGGATGGAAACGCCTAATGATTGAGAAACCGTTCGGGCACGACCTTACATCTGCTCG 473
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                                                                                                      ATTGANCATCTACTACGAAGGAATCNCAGTNAAAATCCTNCAGGTTTTAAGGGTTTCAAAN 222
                                                                      ATTGATCATCTACTANGAANGAATCTCATTTAAAATCTTACAGTTTTAAAGGTTTCAAAT
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                                                                                                                                                       Conservative
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ilarity 55.7%;
Conservative
TTAGCCACTATGAAATCCTACTTACATAGATAA 101
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                                                                                                                                               Score 61.4; DB 17;
Pred. No. 4e-08;
0; Mismatches 33;
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Pred. No. 8e-08;
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                                                                                                                                                 Indels
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                                                                            RESULT 11
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                    Sequence 22036, Application US/10282122A
Publication No. US20040029129A1
GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/09/815,242
CURRENT FILING DATE: 2001-03-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
NAME/KEY: CDS
LOCATION: (1)...(1485)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/269,308 FILING DATE: 2001-02-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 60/253,625
FILING DATE: 2000-11-27
APPLICATION NUMBER: 60/257,931
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FILING DATE: 2000-05-26
APPLICATION NUMBER: 60/242,578
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                                                                                                                                        602
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                                                                                                                                                                                                                                                                                                                               113 GGCTGACAATATCTTCTTTCAAACTTTCAGGAAAAGCAAATATATAGAATTGANCATC 172
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                                                                                                                                      AACCGCTTTGGAACCGTAATTTCATTGATTA 632
                                                                                                                                                                                                                                                    TACTAGGAAGGAATCNCAGTNAAAATCCTNCAGGTTTAAGGGTTTCAAANNNAGNTTTTG 232
                                                                                                                                                                                                                                                                                           CATTAGACGTACAAATCCACCGTTTCTTTGAAGAACACCAAATTTATCGTATCGATCATT
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                                                                                                                                                                          Xu, H. Howard
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Zyskind, Judith W.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yamamoto, Robert T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Grant J.
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Pred. No. 1e-07;
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                                                                                                                                                                                                                                                                                                                             Matches
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PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR FILING DATE: 2000-05-26
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CURRENT APPLICATION NUMBER: US/10/282,122A
CURRENT FILING DATE: 2003-02-20
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Local Similarity 52.1%; Pred. No. 1e-07;
Local Similarity 52.1%; Mismatches 101;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 2001-02-09
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APPLICATION NUMBER: 60/253,625
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                                                                                                                                                                                             GGCTGACACAATATCTTCTTTCAAACTTTCAGGAAAAGCAAATATATAGAATTGANCATC 172
                                                                                                                                                                                                                                                                        AGGGATGGAATCGCATAATATTTGAGAAGCCATTTGGCTTTGATGCACTTTCTTCCCATA 112
                                   AGCCACCTTNGAGNNGNACNTNNNNNGANNA 263
                                                                               Carr, Grant
Yamamoto, Robert
Forsyth, R.
AACCGCTTTGGAACCGTAATTTCATTGATTA
                                                                                                                   TACTAGGAAGGAATCNCAGTNAAAATCCTNCAGGTTTAAGGGTTTCAAANNNAGNTTTTG 232
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Trawick, John
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Haselbeck, Robert
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632
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US-10-329-670-1/c

; Sequence 1, Application US/10329670
; Paublication No. US20040018503A1
; GENERAL INFORMATION:
; APPLICANT: Fleischmann et al.
; TITLE OF INVENTION: Nucleotide Sequence of the Haemophilus influenzae Rd Genome,
; TITLE OF INVENTION: Thereof, and Uses Thereof
; FILE REFERENCE: PB186P1
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CURRENT APPLICATION NUMBER: US/10/329,670
CURRENT FILING DATE: 2002-12-24
PRIOR APPLICATION NUMBER: US 09/643,990
PRIOR FILING DATE: 2000-08-23
PRIOR PRIOR APPLICATION NUMBER: US 08/487,429
PRIOR APPLICATION NUMBER: US 08/487,429
PRIOR APPLICATION NUMBER: US 08/426,787
PRIOR FILING DATE: 1995-04-21
NUMBER OF SEQ ID NOS: 1
SEQ ID NO 1
LENGTH: 1830121
PURDER NEWS OTHER INFORMATION: n equals a, FEATURE:
NAME/KEY: misc_feature
LOCATION: (44416)...(44416)
OTHER INFORMATION: n equals a, FEATURE: NAME/KEY: misc feature LOCATION: (47036)..(47036) COTHER INFORMATION: n equals FEATURE:
NAME/KEY: misc feature
LOCATION: (45732)...(45732)
OTHER INFORMATION: n equals FEATURE:
NAME/KEY: misc_feature
LOCATION: (45593)...(45593)
OTHER INFORMATION: n equals FEATURE:
NAME/KEY: misc_feature
LOCATION: (44975)...(44975)
OTHER INFORMATION: n equals NAME/KEY: misc_feature LOCATION: (51334)..(51334) FEATURE: NAME/KEY: misc feature LOCATION: (44905)..(44905) OTHER INFORMATION: n equals NAME/KEY: misc feature LOCATION: (36551)...(36551) OTHER INFORMATION: n equals NAME/KEY: misc feature LOCATION: (29298) ..(29298) OTHER INFORMATION: n equals LOCATION: (36636)..(36636)
OTHER INFORMATION: n equals LOCATION: (36543)..(36543)
OTHER INFORMATION: n equals NAME/KEY: misc feature LOCATION: (36543)..(36 NAME/KEY: misc feature LOCATION: (10150)..(10150) OTHER INFORMATION: n equals NAME/KEY: misc feature LOCATION: (4747)...(4747) OTHER INFORMATION: n equals JAME/KEY: misc_feature OTHER INFORMATION: n equals NAME/KEY: misc feature LOCATION: (9921)..(9921) ORGANISM: Haemophilus influenzae FEATURE: TYPE: DNA misc_feature (36636)..(36636) a, a, a, a a a a, a, a, 9 a, a) ď, a, ŗ ŗ ŗ , Ţ ŗ ŗ Ĺ Ç 'n ŗ Ĺ , ŗ Q g ιQ g g Q 9 9 g ω Ω ω Q ω õ ç or S, ö 20 å 9 õ õ or 얁 ö Ω a a a a ი ი Ω

NAME/KEY: misc feature LOCATION: (105121)...(105121) OTHER INFORMATION: n equals a, t FEATURE: NAME/KEY: misc feature LOCATION: (107248)...(107248) OTHER INFORMATION: n equals a, t NAME/KEY: LOCATION: NAME/KEY: misc feature LOCATION: (120038)..(120038) OTHER INFORMATION: n equals NAME/KEY: misc feature LOCATION: (122167)...(122167) OTHER INFORMATION: n equals NAME/KEY: misc_feature LOCATION: (121344)..(121344) OTHER INFORMATION: n equals NAME/KEY: misc feature LOCATION: (119750)..(119750) OTHER INFORMATION: n equals FEATURE:
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LOCATION: (117136)..(117136)
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NAME/KEY: misc feature
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NAME/KEY: misc_feature
LOCATION: (51786)...(51786) NAME/KEY: misc feature LOCATION: (65313)..(65313) OTHER INFORMATION: n equals NAME/KEY: misc feature LOCATION: (55369)..(55369) OTHER INFORMATION: n equals NAME/KEY: misc feature LOCATION: (51805)..(51805) OTHER INFORMATION: n equals LOCATION: (65309)..(65309)
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US-10-158-865-1/c
; Sequence 1, Application US/10158865
; Publication No. US20040203093A1
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LOCATION: (152500)..(152500)
OTHER INFORMATION: n equals
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LOCATION: (131360)..(131360)
OTHER INFORMATION: n equals a,
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LOCATION: (152530)..(152530)
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NAME/KEY: misc_feature
*COLTION: (150841)..(150841)
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LOCATION: (147197)..(147197)
OTHER INFORMATION: n equals a,
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LOCATION: (145942)..(145942)
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LOCATION: (145171)...(145171)
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LOCATION: (142750)..(142750)
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LOCATION: (131340)...(131340)
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LOCATION: (140398)..(140398)
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                                                                                      577176
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Similarity 52.1%;
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Pred. No. 3.6e-06;
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SOFTWARE:
SEQ ID NO 1
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APPLICANT: Fleischmann et al.
TITLE OF INVENTION: Thereof, and Uses Thereof
TITLE OF INVENTION: Thereof, and Uses Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 09/557,884
PRIOR FILING DATE: 2000-04-25
PRIOR APPLICATION NUMBER: US 08/476,102
PRIOR FILING DATE: 1995-06-07
PRIOR APPLICATION NUMBER: US 08/426,787
PRIOR APPLICATION NUMBER: US 08/426,787
PRIOR FILING DATE: 1995-04-21
NUMBER OF SEQ ID NOS: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: PB186P2C1D1
CURRENT APPLICATION NUMBER: US/10/158,865
CURRENT FILING DATE: 2002-06-03
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LOCATION: (29298)..(29298)
OTHER INFORMATION: n equals
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LOCATION: (10150)..(10150)
OTHER INFORMATION: n equals
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NAME/KEY: misc_feature
LOCATION: (4747)..(4747)
OTHER_INFORMATION: n equals
                             NAME/KEY: misc feature
LOCATION: (45732)..(45732)
OTHER INFORMATION: n equals
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NAME/KEY: misc_feature
LOCATION: (45593)...(45593)
OTHER_INFORWATION: n equals
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NAME/KEY: misc_feature
LOCATION: (44975)...(44975)
OTHER INFORMATION: n equals
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NAME/KEY: misc_feature
LOCATION: (44905)...(44905)
OTHER INFORMATION: n equals
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LOCATION: (44416)..(44416)
OTHER INFORMATION: n equals
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NAME/KEY: misc_feature
LOCATION: (40808) . . (40810)
OTHER INFORMATION: n equals
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LOCATION: (36551)...(36551)
OTHER INFORMATION: n equals
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LOCATION: (36543)...(36543)
OTHER INFORMATION: n equals
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NAME/KEY: misc_feature
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NAME/KEY: misc_feature
LOCATION: (36636)..(36636)
OTHER INFORMATION: n equals
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NAME/KEY: misc feature
LOCATION: (119924) . (119924)
OTHER INFORMATION: n equals a,
FEATURE:
NAME/KEY: misc feature
LOCATION: (120038) . (120038)
OTHER INFORMATION: n equals a,
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LOCATION: (121344)..(121344)
OTHER_INFORMATION: n equals
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NAME/KEY: misc_feature
LOCATION: (119750)...(119750)
OTHER_INFORMATION: n equals
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LOCATION: (122167)..(122167)
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LOCATION: (117136)...(117136)
OTHER INFORMATION: n equals
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NAME/KEY: misc_feature
LOCATION: (107248)...(107248)
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LOCATION: (105121)..(105121)
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LOCATION: (102696)...(102696)
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LOCATION: (65309)..(65309)
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NAME/KEY: misc_feature
LOCATION: (152530)..(152530)
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LOCATION: (152500)...(152500)
OTHER_INFORMATION: n equals
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LOCATION: (150841)...(150841)
OTHER INFORMATION: n equals
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LOCATION: (147197)...(147197)
OTHER INFORMATION: n equals a,t,
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LOCATION: (145942)..(145942)
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LOCATION: (145058)..(145058)
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NAME/KEY: misc_feature
LOCATION: (145171)..(145171)
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NAME/KEY: misc_feature
LOCATION: (142750)...(142750)
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LOCATION: (131360)..(131360)
OTHER INFORMATION: n equals
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LOCATION: (140398)...(140398)
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NAME/KEY: misc_feature
LOCATION: (131340)...(131340)
OTHER INFORMATION: n equals
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NAME/KEY: misc_feature
LOCATION: (122336)..(122336)
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                                                                                                                                                        577356 ATGGCTGGAAACGTATTATCGTTGAAAAACCTTTTGGTTACGATGAAAAAACAGCGCAAA
                                                      577236
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                         AGCCACCTTNGAGNNGNACNTNNNNNGANNA
                                                   ATTTAGGTAAAGAAACCGTTCAAAACTTGCTCGTATTACGTT
AACCGCTTTGGAACCGTAATTTCATTGATTA 577146
                                                                                                                           GGCTGACACAATATCTTCTTTCAAACTTTTCAGGAAAAGCAAATATATAGAATTGANCATC
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CURRENT APPLICATION NUMBER: US/09/938,842A
CURRENT FILING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: US 60/227,866
PRIOR FILING DATE: 2000-08-24
PRIOR APPLICATION NUMBER: US 60/264,647
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-01-16
PRIOR APPLICATION NUMBER: US 60/300,111
PRIOR APPLICATION NUMBER: US 60/300,111
PRIOR FILING DATE: 2001-06-22
NUMBER OF SEQ ID NOS: 5379
SEQ ID NO 2455
LENGTH: 1782
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; ORGANISM: Arabidopsis thaliana
US-09-938-842A-2455
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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Best Local Similarity
SEQ ID NO 2455
                    APPLICANT: Harper, Jeff
APPLICANT: Kreps, Joel
APPLICANT: Wang, Xun
APPLICANT: Wang, Xun
APPLICANT: Zhu, Tong
TITLE OF INVENTION: STRESS-REGULATED GENES OF PLANTS, TRANSGENIC PLANTS CONTAINING
TITLE OF INVENTION: SAME, AND METHODS OF USE
FILE REFERENCE: SCRIP1300-3
CURRENT APPLICATION NUMBER: US/09/938,842A
CURRENT FILING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: US 60/227,866
PRIOR APPLICATION NUMBER: US 60/264,647
PRIOR APPLICATION NUMBER: US 60/264,647
PRIOR APPLICATION NUMBER: US 60/300,111
PRIOR PILING DATE: 2001-06-22
RUMBER OF SEQ ID NOS: 5379
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Patent No. US20020160378A1
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APPLICANT: Kreps, Joel
APPLICANT: Wang, Xun
APPLICANT: Zhu, Tong
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; ORGANISM: Arabidopsis thaliana US-09-938-842A-2455
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855 GCTAGTCGAGAACTTATCTGTTCTTCGATTCTCAAACCTTATATTTGAGCCGCTATGG 912
                                                                         795 CCTCAAGCAGTÁTTTGGÁGGÁAGATCÁÁATTTTTÁGGÁTAGÁCCÁTTACCTAGGÁAAÁGGÁ 854
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                                   TCNCAGTNAAAATCCTNCAGGTTTAAGGGTTTCAAANNNAGNTTTTGAGCCACCTTNG 243
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Search completed: May 4, 2005, 21:11:33 Job time: 305.387 secs .

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Result
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4 US-09-902-540-4821

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4 US-09-103-840A-2

4 US-09-103-840A-2

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Sequence 1179, Ap
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Sequence 1665, Ap
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Sequence 3533, Ap
Sequence 3533, Ap
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Sequence 213, Appli
Sequence 3106, Ap
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APPLICANT: Lalgudi, Raghunath V.
APPLICANT: Ito, Laura Y.
APPLICANT: Sherman, Bradley K.
TITLE OF INVENTION: POLYNUCLECTIDES AND POLY
FILE REFERENCE: PL-0017 US
CURRENT APPLICATION NUMBER: US/09/313,294A
CURRENT FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 7600
SOFTWARE: PERL Program
SEQ ID NO 6236
LENGTH: 253
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NAME/KEY: misc_feature

OTHER INFORMATION: Incyte ID No. 6476212 700351460H1
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GENERAL INFORMATION:
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Patent No. 6476212
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Patent No. 6476212
GENERAL INFORMATION:
APPLICANT: Lalgudi, Raghunath V.
APPLICANT: Ito, Laura Y.
APPLICANT: Sherman, Bradley K.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN EAR FILE REFERENCE: PL-0017 US
CURRENT APPLICATION NUMBER: US/09/313,294A
CURRENT FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 7600
SEQ ID NO 4756
LENGTH: 277
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LOCATION: 2, 72, 88
OTHER INFORMATION: a, t, c, g,
-09-313-294A-4756
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                                                                                            Score 148; DB 4; 1
Pred. No. 4.9e-31;
0; Mismatches 0;
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US-09-902-540-1237
; Sequence 1237, Application US/09902540
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APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15849)B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 4821
LENGTH: 1404
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Pred. No. 2.3e-19;
0; Mismatches 164;
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GENERAL INFORMATION:
APPLICANT: Goldman,
APPLICANT: Hinkle,

Barry S. Gregory J.

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NAME/KEY: misc_feature
OTHER INFORMATION: Incyte I
NAME/KEY: unsure
LOCATION: 51
OTHER INFORMATION: a, t, c,
S-09-313-294A-5508
                                                                                                                                                                                  SOFTWARE: PER
SEQ ID NO 5508
LENGTH: 294
TYPE: DNA
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Best Local Similarity
Matches 223; Conserv
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PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 1237
LENGTH: 26659
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 6476212
GENERAL INFORMATION:
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APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15849)B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
                                                                                                                                                                                                                                                                                                 FILE REFERENCE: PL-0017 US
CURRENT APPLICATION NUMBER: US/09/313,294A
CURRENT FILING DATE: 1999-05-14
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Lalgudi, Raghunath V. APPLICANT: Ito, Laura Y.
                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Sherman, Bradley K.

TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN EAR
                                                                                                                                         ORGANISM: Zea mays FEATURE:
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                            or other
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US-09-103-840A-2
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Best Local Similarity 88.4
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CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/09103840A Patent No. 6294328
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
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APPLICANT: VENTER, John C.
TITLE OF INVENTION: TUBERCULOSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: FLEISCHMAN,
APPLICANT: WHITE, Ower
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Mycobacterium tuberculosis FEATURE:
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                                      2092028
                                                                                                         2092088
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                                                                                                                                                                                                                                                  2092202 T----TCGTGCGCAGTGAAACGATCCCCGAATTTCTTGCCGCACTGGAAAAACCGCGT
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   GGAGGAAAGGAGGGTGCCCGCCAACGGCCCGTCCTTGATG 410
                                      ATGCGCCGCGAGAAAGCGATGCGTGAGCGGGGCTTGCACTTCGTCGGGGCCGGGATCTCC 2091969
                                                                      GAGAGGAGGAGAAGGCCATGGAGGAGCGCCGCCTNCTGTATCTTGGCATGGGTGTCTCT
                                                                                                         GACGCCATGGAACCCGGCGACATCATCGACGGCGGCAATGCGTTGTACACCGACACC
                                                                                                                                                                             CGGGTGCTGATCATGGTCAAGGCCGGAGAGGCCACTGACGCTGTCATCAACGAACTTGCT
                                                                                                                                                                                                                GTGGTGATCATGCTCGTCAAGGCCGGCGCGCCAGTTGACCCAGACCATCGCGACGCTCGCA
                                                                                                                                                                                                                                                                                   CTTCCCGTCTACGGCTTCCATGACCCCGCGTCCTTTGTGAAGTCCATTCAGAAGCCACGG
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                                                                                                                                                                                                                                                                                                                                                                                                                             GCGGTCATGGGGCAGAACCTTGCCCTCAACATTGCAGAGAAAGGGTTCCCCATCTCTGTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTACAACAGGACAA - CTCCAAGGTTGATGAGACCGTGCAGCGTGCCAAGGTCGAAGGAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CGCGGTCATGGGGCAGAACCTTGCCCTCAACATTGCAGAGAAAGGGTTCCCCCATCTCTGT
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Pred. No. 3.1e-16;
0; Mismatches 174;
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Pred. No. 1.9e-18;
0; Mismatches 15
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3-09-557-884-1/c
5-09-557-884-1/c
Sequence 1, Application US/09557884
PATEDIT NO. 6506581
; GENERAL INFORMATION:
APPLICANT: Fleischmann et al.
APPLICANT: Fleischmann et al.
TITLE OF INVENTION: The Nucleotide sequence of
the Haemophilus influenzae Rd Genome, Fragments
Thereof, and Uses Thereof
                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity Matches 222; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
573896 ATTATTÁCCACÁTTTAGÁAGAGGGCGACATCÁTTATTGÁTGGCGGTAÁCTCAAACTÁTCC 573837
                                                                                                                                                                        574013
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FILING DATE: JUN-5-1995
ATTORNEY/AGENT INFORMATION:
NAME: Michelle S. Marks
REGISTRATION NUMBER: 41,971
REFERENCE/DOCKET NUMBER: PB186P3
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/557,884
FILING DATE: 25-Apr-2000
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3 1/2 inch diskette
COMPUTER: Dell Pentium
COMPUTER: Dell Pentium
OPERATING SYSTEM: MS DOS v6.22
                                                                                                               185 GCCACGGGTGGTGATCATGCTCGAAGGCCGGCGCGCCAGTTGACCAGACCATCGCGAC 244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS: 1
ADDRESSE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
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                                   ACCACGTAAAGTGATGTTAATGGTGCGCGCTGGTGATGTGGTGGATCAATTCATTGAAGC
                                                                                                                                                           AGGCACCAATATTATCGGCGCATACTCTTTAGAAGATTTAGCTGCTAAAT---TAGAAAA
                                                                                                                                                                                                                                       CGTGGCATATAACCGTACCACTTCAAAAGTGGATGAGTTTTTTACAAGGTGC---GGCAAA 574014
                                                                                                                                                                                                                                                                                                                                            CGCGTCCGCGGTCATGGGGCAGAACCTTGCCCTCAACATTGCAGAGAAAAGGGTTCCCCCAT 64
                                                                                                                                                                                               AGGAAACCTTCCCGTCTACGGCTTCCATGACCCCCGCGTCCTTTGTGAAGTCCATTCAGAA 184
                                                                                                                                                                                                                                                                             CTCTGTGTACAACAGGACAACCTCCAAGGTGGACGAGACCGTGCAGGCGTGCCAAGGCAGA 124
                                                                                                                                                                                                                                                                                                                   CGGCTTAĞCCĞTAATĞĞĞĞAAAACCTCATTTTAAATATĞAACĞACCACĞĞĞTTTAAAĞT 574071
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Pred. No. 8.5e-14;
0; Mismatches 178;
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US-09-643-990A-1/c
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Best Local :
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 310-309-8439 INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 08/487,429
FILING DATE: 1995-06-07
APPLICATION NUMBER: 08/426,787
FILING DATE: 1995-04-21
ATTORNEY/AGENT INFORMATION:
NAME: Kenley K. Hoover
REGISTRATION NUMBER: 40,302
REFERENCE/DOCKET NUMBER: PB186P10
TELEPONE: 301-610-5790
                                                                                                                      574130
                                                                                                                                                                                                           Local
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APPLICATION NUMBER: US/09/643,990A
FILING DATE: 23-Aug-2000
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3 1/2 inch diskette
COMPUTER: Dell Pentium
OPERATING SYSTEM: MS DOS v6.22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genom
STREET: 9410 Key West
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Robert D. Fleischmann
Mark D. Adams
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                                                                                                                                            5 CGCGTCCGCGTCATGGGGCAGAACCTTGCCCTCAACATTGCCAGAGAAAAGGGTTCCCCAT 64
                                                                                                                                                                                                       h 21.9%;
Similarity 54.7%;
                                  CGTGGCATATAACCGTACCACTTCAAAAGTGGATGAGTTTTTACAAGGTGC---GGCAAA 574014
                                                                                                           CGGCTTAGCCGTAATGGGGCAAAACCTCATTTTAAATATGGAACGACCACGGGTTTAAAGT 574071
AGGAAACCTTCCCGTCTACGGCTTCCATGACCCCGCGTCCTTTGTGAAGTCCATTCAGAA 184
                                                                       CTCTGTGTACAACAGGACAACCTCCAAGGTGGACGAGACCGTGCAGGCGTGCCAAGGCAGA 124
                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
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COUNTRY: U
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Owen White
Hamilton O. Smith
J. Craig Venter
the Hacleotide sequence of
the Haemophilus influenzae
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                                                                                                                                                                                                       Score 89.8; DB 4;
Pred. No. 8.5e-14;
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RESULT 10
US-09-198-452A-1/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence TITLE OF INVENTION: thereof and uses thereof, in particul TITLE OF INVENTION: and treatment of infection FILE REFERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Griffais,
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TYPE: DNA
                     NAME/KEY: misc_feature
LOCATION: (135001)..(15000)
LOCATION: n=a or c or
NAME/KEY: misc_feature
LOCATION: (150001)..(165000)
                                                                                                                            NAME/KEY: misc feature
LOCATION: (120001)...(135000)
OTHER INFORMATION: n=a or c or
                                                                                                                                                                                               LOCATION: (90001)..(105000)
OTHER INFORMATION: n=a or c or g or
NAMB/KEY: misc_feature
LOCATION: (105001)..(120000)
OTHER INFORMATION: n=a or c or g or
NAME/KEY: OTHER INFORMATION: n=a or c or g or
                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
LOCATION: (75001)..(90000)
OTHER INFORMATION: n=a or c
NAME/KEY: misc_feature
LOCATION: (90001)..(105000)
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LOCATION: (60001)..(75000)
OTHER INFORMATION: n=a or c
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OTHER INFORMATION: n=a or c
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LOCATION: (30001)..(45000)
COTHER INFORMATION: n=a or c
NAME/KEY: misc_feature
LOCATION: (45001)..(60000)
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OTHER INFORMATION: n=a or c
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HER INFORMATION: n=a or c or g
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LOCATION: (480001) .. (495000)
OTHER INFORMATION: n=a or c o
NAME/KEY: misc feature
LOCATION: (495001) .. (510000)
OTHER INFORMATION: n=a or c o
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OTHER INFORMATION: n=a or c o
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LOCATION: (40500) . (420000)
OTHER INFORMATION: n=a or c o
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OTHER INFORMATION: n=a or c o)
NAME/KEY: misc feature
LOCATION: (315001)..(330000)
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NAME/KEY: misc_feature
LOCATION: (510001) .. (525000)
OTHER INFORMATION: n=a or c
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LOCATION: (465001)..(480000)
OTHER INFORMATION: n=a or c
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LOCATION: (375001) ...(390000)
OTHER INFORMATION: n=a or c
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LOCATION: (330001)..(345000)
OTHER INFORMATION: n=a or c
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LOCATION: (28501)..(30000)
OTHER INFORMATION: n=a or c
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LOCATION: (435001)..(450000)
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NAME/KEY: misc feature
LOCATION: (360001)..(375000)
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LOCATION: (25501)..(270000)
OTHER INFORMATION: n=a or c
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LOCATION: (225001) . (240000)
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LOCATION: (195001)...
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LOCATION: (70501)...(720000)
OTHER INFORMATION: nea or c or g or
NAME/KEY: misc feature
LOCATION: (720001)...(735000)
OTHER INFORMATION: nea or c or g or
NAME/KEY: misc feature
LOCATION: (720001)...(750000)
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NAME/KEY: misc feature
LOCATION: (755001)...(750000)
OTHER INFORMATION: nea or c or g or
NAME/KEY: misc feature
LOCATION: (75001)...(765000)
OTHER INFORMATION: nea or c or g or
NAME/KEY: misc feature
LOCATION: (755001)...(780000)
OTHER INFORMATION: nea or c or g or
NAME/KEY: misc feature
LOCATION: (75001)...(795000)
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NAME/KEY: misc feature
LOCATION: (780001)...(795000)
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LOCATION: (79501).. (81000)
OTHER INFORMATION: n=a or c or
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LOCATION: (81001).. (82500)
OTHER INFORMATION: n=a or c or
NAME/KEY: misc feature
LOCATION: (825001).. (84000)
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NAME/KEY: misc feature
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LOCATION: (645001)..(660000)
OTHER INFORMATION: n=a or c or g or
NAME/KEY: misc_feature
LOCATION: (660001)..(675000)
OTHER INFORMATION: n=a or c or g or
NAME/KEY: misc_feature
LOCATION: (675001)..(690000)
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NAME/KEY: misc_feature
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LOCATION: (540001) .. (555000)
CTHER INFORMATION: n=a or c or g
NAME/KEY: misc feature
LOCATION: (555001) .. (570000)
CTHER INFORMATION: n=a or c or g
LOCATION: (555001) .. (585000)
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LOCATION: (570001) .. (585000)
CTHER INFORMATION: n=a or c or g
NAME/KEY: misc feature
LOCATION: (585001) .. (600000)
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LOCATION: (600001) .. (615000)
CTHER INFORMATION: n=a or c or g
NAME/KEY: misc feature
LOCATION: (615001) .. (630000)
CTHER INFORMATION: n=a or c or g
NAME/KEY: misc feature
LOCATION: (615001) .. (630000)
CTHER INFORMATION: n=a or c or g
NAME/KEY: misc feature
LOCATION: (630001) .. (645000)
CTHER INFORMATION: n=a or c or g
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                                                                                                                                     Query Match
Best Local S
Matches 212
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SOPTWARE: FastSEQ for Windows Version
SEQ ID NO 1
LENGTH: 1230230
TYPE: DNA
ORGANISM: Chlamydia pneumoniae
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APPLICANT: Stephens, Richard
APPLICANT: Mitchell, Wayne
APPLICANT: Kalman, Sue
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Best Local S
Matches 213
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APPLICANT: Kalman, Sue
APPLICANT: Lavis, Ronald
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Chlamydia Pneumoniae Genome Sequence
FILE REFERENCE: 018941-000411US
CURRENT APPLICATION NUMBER: US/09/438,185A
CURRENT FILING DATE: 2002-03-13
                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/108,279
PRIOR FILING DATE: 1998-11-12
PRIOR APPLICATION NUMBER: US 60/128,606
PRIOR FILING DATE: 1999-04-08
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                                                       405328 GCTGTCATGGGGAAAAATCTTGTCTTAAACATGATAGATCATGGTTTTTCTGTCTCTGTC 405269
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72 TACAACAGGACAACCTCCAAGGTGGACGAGACCGTGCAGCGTGCCAAGGCAGAAGGAAAC 131
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                                                                                           GCGGTCATGGGGCAGAACCTTGCCCTCAACATTGCAGAGAAAGGGGTTCCCCCATCTCTGTG 71
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                                                                                                                             Score 88; DB 4; Length 123
Pred. No. 2.3e-13;
0; Mismatches 181; Indels
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Pred. No. 8.4e-14;
); Mismatches 180;
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APPLICANT: FRASER, Claire M.
APPLICANT: VENTER, John C.
TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN
TITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TENGTH: 4411529
TYPE: DNA
ORGANISM: Mycobacterium tuberculosis
OTHER INFORMATION: H37Rv
  2094835
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hes 218;
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                                                                                                                                                                               GTGCTGATCATGGTCAAGGCCGGAGAGGCCACTGACGCTGACGCTGTCATCAACGAACTT 2094896
                                                                                                                                                                                                                                                                     TT---CGTGCGCAGTGAAACGATCCCCGAATTTCTTGCCGCACTGGAAAAACCGCGTCGG
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  ACCATGCGCCGCGAGAAAGCGATGCGTGAGCGGGGCTTGCACTTCGTCGGGGGCCGGGATC
                                          ACGGAGAGGAGGAAAGGCCATGGAGGAGCGCGGCCTNCTGTATCTTGGCATGGGTGTC 368
                                                                                      GCTGACGCCATGGAACCCGGCGACATCATCATCGACGGCGGCAATGCGTTGTACACCGAC
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Pred. No. 1.4e-12;
0; Mismatches 178;
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US-09-543-681A-1208
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CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 1179
                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: GARY BRETON
                                                                                                                                                                                              Sequence 1208, App
Patent No. 6605709
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Best Local
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Patent No. 661083
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: 2709.1002-001
CURRENT APPLICATION NUMBER: US/09/543,681A
CURRENT FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 60/128,706
PRIOR PILING DATE: 1999-04-09
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
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ORGANISM: Klebsiella
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCTTACCTTGATAAAGGCGACATCATGATGATGGCGGCAACACCTTCTTCCAGGACACT 390
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                                                                                                                                                                                                                   Application US/09543681A
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Pred. No. 5.5e-11;
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                                                                                                                                                                                                                                                                                                                                                                    410
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NUMBER OF SEQ ID NOS: 8344

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                                                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: synthetic
; OTHER INFORMATION: nucleic acid sequence
US-09-710-279-1665
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US-09-710-279-1665
; Sequence 1665, Application US/09710279
; Patent No. 6703492
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; ORGANISM: Proteus mirabilis
US-09-543-681A-1208
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                                                                                                                               Query Match
Best Local S
Matches 197
                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: KIMMERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1665
LENGTH: 1407
TYPEE. DAY
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LENGTH: 1461
TYPE: DNA
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Best Local Similarity 49.9%;
Matches 199; Conservative
                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                              TYPE: DNA
                                                                                                                                 Local Similarity
les 197; Conserv
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TATAACCGATCAAGACAAAAAACTGATGAAATGGTTAAAGAATCGCCTGGAAGAGAAATT 150
                            ТАСААСАGGACAACCTCCAAGGTGGACGAGACCGTGCAGGCGTGCCAAGGCAGAAGGAAAC 131
                                                            GCAGTAATGGGGAAAAACCTAGCTTGGAATATTGAATCACGTGGTTATAGTGTTTCTGTT
                                                                                          ecegrearegecaeaaccrrecceaacarrecaeasaaaaeegrreccearcrere
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GCTCACTTGGAGCAGGGCGACTGCATCATCGATGGGGGAACGAGTGGTACGAGAACACG 311
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                                                                                                                                 Conservative
                                                                                                                          15.3%; Score 62.8; DB 4; Length 1 49.4%; Pred. No. 1.7e-07; tive 0; Mismatches 193; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                          Length 1407;
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                          egaegaaaegereccecaacegeccercerreare
                                                                                   GAGAGGAGGAGAAGGCCATGGAGGAGCGCCGCCTNCTGTATCTTGGCATGGGTGTCTCT 371
                                                                                                                                                                          AAGATTTTATTAATGGTAAAAGCTGGACCTGCAACAGATGCCACTATAGATGGTTTATTA 261
                                                                                                                                                                                                                                  TACCCAACATACTCATTAGAAGA-----ATTTGTAGAATCTTTAGAGAAACCTCGT 201
                                                                                                                  CCTTTATTAGACGATGATATTTTAATTGATGGTGGTAATACTAATTACCAAGATACG
                                                                                                                                            GCTCACTTGGAGCAGGGGACTGCATCATCGATGGGGGGAACGAGTGGTACGAGAACACG 311
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420
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Search completed: May 4, 2005, 12:27:54 Job time : 137.974 secs

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Minimum DB seq length: 0
Maximum DB seq length: 200000000
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scoring table:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Title:
Perfect score:
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1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*

3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*

5: /cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*

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7: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
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Gapop 10.0 , Gapext 1.0
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Copyright (c) 1993 - 2005 Compugen Ltd
/cgn2_6/ptodata/1/pubpna/US09E PUBCOMB.seq:*
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/cgn2_6/ptodata/1/pubpna/US10 NEW PUB.seq:*
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/cgn2_6/ptodata/1/pubpna/US10 NEW PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10 NEW PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

11	• •	8	7	6	υ	.4.	ω	2	_	Result No.
396.4	396.4	396.4	396.4	396.4	396.4	396.4	396.4	396.4	396.4	Score
96.7	96.7	96.7	96.7	96.7	96.7	96.7	96.7	96.7	96.7	Query Match
1691	1674	1672	1672	1671	1669	1661	1624	1587	748	Query Match Length DB
17	17	17	17	17	17	17	17	17	18	BB
US-10-425-114-28011	US-10-425-114-13683	US-10-425-114-32227	US-10-425-114-24039	US-10-425-114-1523	US-10-425-114-31534	US-10-425-114-24375	US-10-425-114-30841	US-10-425-114-6537	US-10-425-115-91003	ID
Sequence 28011, A	Sequence 13683, A			Sequence 1523, Ap	Sequence 31534, A	Sequence 24375, A	Sequence 30841, A	Sequence 6537, Ap	Sequence 91003, A	Description

4.5	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12
207.6	л	230	31	234.8	42	42	262	2	22	323.8	23	23	326	346.8	360	σ	91	93.	93.	93.	93.	ü	93.	93.	93.	93.	93.	93.	94.	94.	96.	96.	9
50.6			•	•	•	•	•	•		•	٠	•	•	•	•	•	•	•	•		•	٠	•	•	•		•	•	٠	•	•	•	•
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S-10-424-599-660	09-294-093B-522	-10-424-599-599	-10-425-114-	-10-424-599-	-10-424-599-	-10-425-11	-10-739-930-	5-115-	-10-425-115-	5-115-12	-10-425-114-3	-10-425-114-3	-10-437-963-9	0-767-701	0-425-115	-10-425-114	US-10-425-114-26662	-10-425-114	0-425-11	-10-425-11	-10-425-114-3	-10-425-11	-10-425-114-4	0-425-114-2503	-10-425-114-2	-10-425-114-1353	4-4	-10-425-114-3	0-425-115-9	-10-425-114-:	-10-425-115-	-425-114-5	-10-425-114-
e 66028	5229,	e 59924,	equence 12849,	e 59918,	e 76904,	e 10641,	equence 1129,	equence 54519,	e 4492	e 127057	e 3640,	e 32383,	e 98959,	e 14931,	e 91011,	e 12436,	æ	e 1418,	e 24479,	e 4434,	e 3615,	e 25153,	e 4096,	e 25038,	e 24256,	e 13538,	e 4238,	e 3149,	e 91002,	e 2633,	e 91000,	e 508,	e 3294,

ALIGNMENTS

```
APPLICANT: Kovalic, David K.

APPLICANT: Kovalic, David K.

APPLICANT: Zhou, Yihua

APPLICANT: Cao, Yongwei

ITITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

ITITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

ITITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

ITITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

ITITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

ITITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

ITITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

ITITLE OF INVENTION: 2203-04-28

INVENTION NUMBER: US/10/425,115

INVENTION NUMBER: US/30/425,115

INVENTION NUMBER:
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US-10-425-115-91003
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Publication No. US20040214272A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                              397;
                                                                                                                                                                                                                       205 GCGGTCATGGGGCAGAACCTTGCCCTCAACATTGCAGAGAAAGGGTTCCCCCATCTCTGTG
265 TACAACAGGACAACCTCCAAGGTGGACGAGACCGTGCAGCGTGCCAAGGCAGAAGGAAAC
                                                                           72 TACAACAGGACAACCTCCAAGGTGGACGAGACCGTGCAGCGTGCCAAGGCAGAAGGAAAC 131
                                                                                                                                                                                                                                                                                 12 GCGGTCATGGGGCAGAACCTTGCCCTCAACATTGCAGAGAAAGGGTTCCCCCATCTCTGTG 71
                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                         96.7%; Score 396.4; DB 18; Length 748; 99.5%; Pred. No. 2.7e-113; rative 0; Mismatches 2; Indels 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
           324
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LELICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE REFERENCE: 38-21(5313)B
CURRENT FILING DATE: 2013-04-28
NUMBER OF SEG ID NOS: 73128
SEQ ID NO 6537
LENGTH: 1587
TYPE: DNA
CORGANISM: Zea mays
FEATURE:
COTHER INFO--
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Best Local Similarity
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GAGAGGAGGAGAAGGCCATGGAAGAGCGCGCCTNCTGTATCTTGGCATGGGTGTCTCT
                                                                  GCTCACTTGGAGCAGGGCGACTGCATCATCGATGGGGGGAACGAGTGGTACGAGAACACG
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                                              GCTCACTTGGAGCAGGGGGACTGCATCATCGATGGGGGGAACGAGTGGTACGAGAACACG
                                                                                                                         GTGGTGATCATGCTCGTCAAGGCCGGCGGCGACGTTGACCAGACCATCGCGACGCTCGCA
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                                                                                                                                                                                 CTTCCCGTCTACGGCTTCCATGACCCCGCGTCCTTTGTGAAGTCCATTCAGAAGCCACGG
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99.5%;
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Pred. No. 3e-113;
0; Mismatches 2;
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; OTHER INFORMATION: Clone ID: UC-ZMFLB73088H07_FLI
US-10-425-114-30841
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-425-114-30841
; Sequence 30841, Application US/10425114
; Publication No. US20040034888A1
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APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Green, Steven B
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REPERENCE: 38-21(53313)B
CURRENT APPLICATION UNMEER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 30841
LENGTH: 1624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
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                                GGAGGAAAGGAGGGTGCCCGCAACGGCCCGTCCTTGATG 410
                                                                                     GAGAGGAAGGAAGGCCATGGAGGAGCGCGCCTNCTGTATCTTGGCATGGGTGTCTCT
                                                                                                                                                          GCTCACTTGGAGCAGGGCGACTGCATCATCGATGGGGGGAACGAGTGGTACGAGAACACG
                                                               GAGAGGAGGAGAAGGCCATGGAGGAGCGCGGCCTCCTGTATCTTGGCATGGGTGTCTCT
                                                                                                                                   GCTCACTTGGAGCAGGGGACTGCATCATCGATGGGGGGAACGAGTGGTACGAGAACACG
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RESULT 4
US-10-425-114-24375
US-10-425-114-24375
Sequence 24375, Application US/10425114
Publication No. US20040034888A1
GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.

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RESULT 5
US-10-425-114-31534
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APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT APPLICATION NUMBER: US/10/425,114
SEQ ID NO 24375
LENGTH: 1661
                                                                               APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 31534
LENGTH: 1669
TYPEF. NAN
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 31534, Application US/10425114
Publication No. US20040034888A1
                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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                                       TYPE: DNA
ORGANISM: Zea mays
  OTHER INFORMATION: Clone ID: UC-ZMFLB73186B12_FLI
                         FEATURE:
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nes 397; Conservative
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APPLICANT: Zhou, Yihua

APPLICANT: Kovalic, David K.

APPLICANT: Tabaska, Jack E

APPLICANT: Tabaska, Jack E

APPLICANT: Tabaska, Jack E

APPLICANT: Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

FILE REFERENCE: 38-21(53313)B

CURRENT APPLICATION NUMBER: US/10/425,114

CURRENT TAPPLICATION NUMBER: US/10/425,114

NUMBER OF SEQ ID NOS: 73128

SEQ ID NO 1523

LENGTH: 1671

TYPE: DNA

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Best Local
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                                                                                                                                                                                             Similarity
CTTCCCGTCTACGGCTTCCATGACCCCGCGTCCTTTGTGAAGTCCATTCAGAAGCCACGG 191
                                                         TACAACAGGACAACCTCCAAGGTGGACGAGACCGTGCAGGGTGCCAAGGCAGAAGGAAAG 131
                                                                                                         GCGGTCATGGGGCAGAACCTTGCCCTCAACATTGCAGAGAAAGGGTTCCCCCATCTCTGTG 176
                                                                                                                                          GCGGTCATGGGGCAGAACCTTGCCCTCAACATTGCAGAGAAAGGGTTCCCCATCTCTGTG 71
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                                    TACAACAGGACAACCTCCAAGGTGGACGAGACCGTGCAGCGTGCCAAGGCAGAAGGAAAC 236
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                                                                                                                                                                                           96.7%;
99.5%;
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                                                                                                                                                                             Score 396.4; DB 17;
Pred. No. 3e-113;
0; Mismatches 2;
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Pred. No. 3e-113;
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                                                                                                                                                                                                              Length 1671;
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Sequence 24039, Application US/10425114

Publication No. US20040034888A1

GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Screen, Steven E
APPLICANT: Soreen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Gao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With.
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TITLE OF INVENTION: Nucleic Acid Molecules Associ
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; OTHER INFORMATION: Clone ID: LIB3607-006-H2_FLI
US-10-425-114-24039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 7
US-10-425-114-24039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches 397; Conservative
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                                                                                GAGAGGAAGGAAGAGATGAAGGAGGAGGAGCGCCTNCTGTATCTTGGCATGGGTGTCTCT 371
                                                GAGAGGAĞĞAĞAĞĞAĞĞCĞATĞĞAĞĞĞĞĞĞĞĞĞĞĞĞĞTĞTATÇTTĞĞĞATĞĞĞTĞTĞTÇTÇT
                                                                                                                                                                                 GCTCACTTGGAGCAGGGCGACTGCATCCATCGATGGGGGGAACGAGTGGTACGAGAACACG
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                                                                                                                                                GCTCACTTGGAGCAGGGGGACTGCATCATCGATGGGGGAACGAGTGGTACGAGAACACG
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RESULT 9
US-10-425-114-13683
(Sequence 13683, Application US/10425114)
(Publication No. US20040034888A1)
(GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Shou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
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US-10-425-114-32227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David
APPLICANT: Screen, Steven
APPLICANT: Tabaska, Jack
APPLICANT: Cao, Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches 397;
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Best Local Similarity
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Publication No. US20040034888A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Zea mays
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                                                                                                                                                                                                                                                                                                                         478
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; OTHER INFORMATION: Clone ID: UC-ZMFLB73276H04_FLI US-10-425-114-32227
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APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT APPLICATION SUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 32227
LENGTH: 1672
                                                                                                                                                                                                                         252 GCTCACTTGGAGCAGGGCGACTGCATCATCGATGGGGGGAACGAGTGGTACGAGAACACG
                                                                                                                                                                                                                                                                                           298 GTGGTGATCATGCTCGTCAAGGCCGGCGCGCCAGTTGACCAGACCATCGCGACGCTCGCA
                          GGAGGAAAGGAGGGTGCCCGCAACGGCCCGTCCTTGATG 410
                                                                                                                           GAGAGGAGGAGGAAGGCCATGGAGGAGGAGCGCCTNCTGTATCTTGGCATGGGTGTCTCT
                                                                                           GAGAGGAGGAAGGCCATGGAGGAGCGCGCCTCCTGTATCTTGGCATGGGTGTCTCT
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Pred. No. 3e-113;
0; Mismatches 2; Indels 0;
516
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US-10-425-114-446
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; OTHER INFORMATION: Clone ID: US-10-425-114-446
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                                                                                                         NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 446
LENGTH: 1675
                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 446, Application US/10425114
Publication No. US20040034888A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 99.9
Matches 397; Conservative
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LENGTH: 1674
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                                                                                                                                                                     APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cab, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
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TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERNCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
                                                                                                                                                                                                                                                                                                      APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
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                                         ORGANISM: Zea mays FEATURE:
                                                                                     TYPE: DNA
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99.5%;
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Pred. No. 3e-113;
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FILE REFERENCE: 38-21 (53313) B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 28011
LENGTH: 1691
TYPE: DNA
ORGANISM: Zea mays
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; OTHER INFORMATION: Clone ID: LIB4743-047-F5_FLI
US-10-425-114-28011
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US-10-425-114-28011
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APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David
APPLICANT: Screen, Steven
APPLICANT: Tabaska, Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 28011, Application US/10425114 Publication No. US20040034888A1 GENERAL INFORMATION:
                                                                                                                                                                                                                         Query Match
Best Local Similarity
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TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
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                   CTTCCCGTCTACGGCTTCCATGACCCCGCGTCCTTTGTGAAGTCCATTCAGAAGCCACGG 191
                                                                                      TACAACAGGACAACCTCCAAGGTGGACGAGACCGTGCAGCGTGCCAAGGCAGAAGGAAAC 131
                                                                                                                                      GCGGTCATGGGGCAGAACCTTGCCCTCAACATTGCAGAGAAAGGGTTCCCCCATCTCTGTG 193
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                                                                    TACAACAGGACAACCTCCAAGGTGGACGAGACCGTGCAGCGTGCCAAGGCAGAAGGAAAGC 253
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Screen, Steven E
Tabaska, Jack E
                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                         Score 396.4; DB
Pred. No. 3e-113;
0; Mismatches
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Pred. No. 3e-113;
0; Mismatches
                                                                                                                                                                                                                                         DB 17;
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Sequence 3394, Application US/10425114

Publication No. US20040034888A1

GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Chou, Yihua
APPLICANT: Screen, Steven E
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)8

FILE REFERENCE: 38-21(53313)8

CURRENT APPLICATION NUMBER: US/10/425,114

CURRENT FILING DATE: 2003-04-28

SEQ ID NO 3294

LENGTH: 1781
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US-10-425-114-3294
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Best Local
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ORGANISM: Zea mays
FEATURE:
                                                                                                                                                                  407
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397; Conserv
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GGAGGAAAGGAGGGTGCCCGCAACGGCCCGTCCTTGATG 410
                                          GAGAGGAĞĞĞAĞAAĞĞCCATĞĞAĞĞAĞCĞCĞĞCCTCCTĞTATCTTĞĞCATĞĞĞTĞTCTCT
                                                                GAGAGGAGGAGGAGAGCCATGGAGGAGCGCCGGCCTNCTGTATCTTGGCATGGGTGTCTCT
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Pred. No. 3.1e-113;
0; Mismatches 2;
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Sequence 508, Application US/10425114

Publication No. US20040034888A1

GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Cou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 39-21(53313) B
CURRENT APPLICATION NUMBER: US/10/425,114

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 73128

EBQ ID NO 508
LENGTH: 1798
Sequence 91000, Application US/10425115
Publication No. US20040214272A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
TITILE OF INVENTION: Plants
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US-10-425-114-508
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ORGANISM: Zea mays
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Pred. No. 3.1e-113;
0; Mismatches 2;
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ION: Plants 38-21(53222)B

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RESULT 15
US-10-425-114-2633
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                                                                                        ; OTHER INFORMATION: Clone ID: 700219903_FLI US-10-425-114-2633
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                                                                                                                                                                                         APPLICANT: Zhou, Yihua APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated I
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 2633
LENGTH: 2171
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publication No. US20040034888A1
GENERAL INFORMATIN:
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CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 91000
LENGTH: 3520
Query Match 96.3%; Score 394.8; DB 17; Length 2171; Best Local Similarity 99.2%; Pred. No. 1e-112; Matches 396; Conservative 0; Mismatches 3; Indels 0;
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Best Local 9
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APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, Davi
                                                                                                                                TYPE: DNA
ORGANISM: Zea mays
FEATURE:
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ORGANISM: Zea mays
FEATURE:
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Local Similarity 99.5%;
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Search completed: May 4, 2005, 21:11:35 Job time: 467.167 secs THIS PAGE BLANK (USPTO)

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Minimum
Maximum
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Match
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1: /cgm2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgm2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgm2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgm2_6/ptodata/1/ina/6B_COMB.seq:*

5: /cgm2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgm2_6/ptodata/1/ina/backfiles1.seq:*
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US-09-710-279-185
US-09-710-279-371
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US-09-710-279-371
US-09-949-016-17375
US-08-781-986A-454
US-09-949-016-168377
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              Sequence 3185, Ap
Sequence 2798, App
Sequence 185, App
Sequence 3371, Ap
Sequence 17976, Ap
Sequence 17976, Ap
Sequence 65898, App
Sequence 65898, App
Sequence 11, Appli
Sequence 16923, A
Sequence 16837, A
Sequence 11837, A
Sequence 14856, A
Sequence 1485, A
Sequence 1485, A
Sequence 544, Appli
Sequence 544, Appli
Sequence 544, Appli
Sequence 547, Appli
Sequence 549, Appli
Sequence 31, Appli
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Sequence 53690, A
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p	RESUI US-02: See : See : See : See : Paal : Hall :	000
2 TITTTGCAGTTAGTAGAATA	WESULT 1 US-09-248-796A-3185 (Sequence 3185, Application US/09248796A) Patent No. 6747137 GENERAL INFORMATION: APPLICANT: Keith Weinstock et al TITLE OF INVENTION: WICLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: 107196.132 CURRENT APPLICATION NUMBER: US/09/248,796A CURRENT PILING DATE: 1999-02-12 PRIOR APPLICATION NUMBER: US 60/074,725 PRIOR APPLICATION NUMBER: US 60/096,409 PRIOR APPLICATION NUMBER: US 60/096,409 PRIOR FILING DATE: 1998-08-13 NUMBER OF SEQ ID NO 3185 SEQ ID NO 3185 LENGTH: 744 TYPE: DNA CORGANISM: Candida albicans US-09-248-796A-3185 Ouery Match Best Local Similarity 53.8%; Pred. No. 8.9e-07; Matches 128; Conservative 0; Mismatches 104; Indels 6; Gaps 1;	C 28 29.4 12.6 1533 4 US-09-270-767-12691 Sequence 12691, A 29 29.4 12.6 14335 4 US-09-596-002-11 Sequence 11, Appl 29.4 12.6 24405 4 US-09-949-016-12346 Sequence 12040, A 29.4 12.6 250715 4 US-09-949-016-13294 Sequence 13294, A 29.4 12.6 393753 4 US-09-949-016-14573 Sequence 13294, A 29.4 12.6 393753 4 US-09-949-016-14574 Sequence 14574, A 29.4 12.6 818128 4 US-09-949-016-14546 Sequence 14546, A 29.4 12.6 818128 4 US-09-949-016-14547 Sequence 14546, A 29.4 12.6 818128 4 US-09-949-016-14549 Sequence 14547, A 29.4 12.6 818128 4 US-09-949-016-14549 Sequence 14549, A 29.4 12.6 818128 4 US-09-949-016-14550 Sequence 14550, A 29.4 12.6 818128 4 US-09-949-016-14551 Sequence 14552, A 29.4 12.6 818128 4 US-09-949-016-14551 Sequence 14552, A 29.4 12.6 818128 4 US-09-949-016-14553 Sequence 14553, A 29.4 12.6 818128 4 US-09-949-016-14553 Sequence 14553, A 29.4 12.6 818128 4 US-09-949-016-14553 Sequence 14555, A 29.4 12.6 818128 4 US-09-949-016-14555 Sequence 14555, A 29.4 12.6 818128 4 US-09-949-016-14556 Sequence 14556, A 29.4 12.6 818128 Sequence 14556, A 29.4 12.6 81812

RESULT 2
US-09-134-001C-2798
; Sequence 2798, Application US/09134001C
; Patent No. 6380370
; Patent No. MATION:
; GENERAL INFORMATION:
; APPLICANT: Lynn Doucette-Stamm et al
; APPLICANT: Lynn Doucette-Stamm et al

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RESULT 4
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CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR FILING DATE: 1997-11-08
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674
SEQ ID NO 2798
SEQ ID NO 2798
LENGTH: 507
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; ORGANISM: Staphylococcus epidermidis
US-09-134-001C-2798
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                                                                           Sequence 3371, Application US/09710279 Patent No. 6703492 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 185
                                                                                                                                                                                                                                                                                                                                             Matches
       APPLICANT: KIMMERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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CURRENT APPLICATION NUMBER: US/09/710,279
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Best Local
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TITLE OF INVENTION: STARHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS
FILE REFERENCE: PU34880US
CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: nucleic acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence
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                                                                         INFORMATION:
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                                                                                                                                                                                                                                                               TTAGAAGATTTAATTAATGACCACGAAGATACTGGGCAATTATTAAATGTTATGAATCAA 495
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Pred. No. 0.1;
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Pred. No. 0.095;
0; Mismatches 34;
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                                                                                                                                                        US-09-949-016-17375/c
Sequence 17375, Application US/09949016
PATENT NO. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHIEMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-710-279-3794
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: KIMMERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
CURRENT APPLICATION NUMBER: U5/09/710,279
CURRENT FILING DATE: 2000-11-09
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR PILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFTWARE: PSECNII VEY: 2.1
SEQ ID NO 3794
                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
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SEQ ID NO 3371
LENGTH: 2945
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Best Local (
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OTHER INFORMATION: Description of Artificial Sequence: synthetic
OTHER INFORMATION: nucleic acid sequence
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PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
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Local Similarity 61.8%;
                                                                                                                                                                                                                                                                                                 780 TTAGAAGATTTAATTAATGACCACGAAGATACTGGGCAATTATTAAATGTTATGAATCAA
                                                                                                                                                                                                                                720 CTAACCTCTGATTATCAAACCCCAGAAGA 692
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                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                       ASSOCIATED
OF DETECTI
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                                       DETECTION
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                                   AND USES THEREOF
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US-08-956-171E-454
Sequence 454, Application US/08956171E
Patent No. 6593114
GENERAL INFORMATION:
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Best Local Similarity
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PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 4.0 EQ ID NO 17375
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NAME/KEY: misc_feature
LOCATION: (1)...(84296)
OTHER INFORMATION: n = A,T,C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Craig A. Rosen
Steven C. Barash
Michael R. Fannon
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
NUMBER OF SEQUENCES: 5256
                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
ATTORNEY/AGENT INFORMATION: NAME: Mark J. Hyman
                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      172 GATGATAAGATCATATTGCATG 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          112 AAGCTACCGCAGATTATTTTATTTGGCATTGCCTCCATCAGTCTACCCATCAGTATGCGA 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              52 ATTTGAGTTATTGAATGAGGCAATCTCTGAGTATGAGACTTCAGAAAACAATGACTCGGG 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        73;
                                 APPLICATION NUMBER: 60/009,861
FILING DATE: January 5, 1996
APPLICATION NUMBER: 08/781,986
FILING DATE: January 3, 1997
                                                                                                                                           APPLICATION NUMBER: US/08/956,171E
FILING DATE: 20-Oct-1997
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                 COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
                                                                                                                                                                                                                                                                                                                                                                                   STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 9410 Key West Avenue
                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
                                                                                                                           APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Human Genome Sciences,
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Gil H. Choi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patrick S. Dillon
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Pred. No. 10;
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US-08-781-986A-454
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                                                                                                                           NAME: Benson, Bob
REGISTRATION NUMBER: 30,446
REGISTRATION NUMBER: PB24
REFERENCE/DOCKET NUMBER: PB24
TELECOMUNICATION INFORMATION:
TET.EPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 454: SEQUENCE CHARACTERISTICS:
                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 4358 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: ASCII Text CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: HP Vecu
                                   STRANDEDNESS:
                                                        TYPE:
                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Rockville
STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 13.4%;
Local Similarity 53.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      168 GCGA 171
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               48 AAGGATTTGAGTTATTGAATGAGGCAATCTCTGAGTATGAGACTTCAGAAAAACAATGACT 107
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                                                    nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 4358 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (240) 314-1224
TELEFAX: (301) 309-8439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: PB248P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9410 Key West Avenue
                                                                     4358 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                    linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Diskette,
                                   double
                                                                                                                         309-8512
                                                                                                                                                                                                                                                                                                                                                                                                 MSDOS version 6.2
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                                                                                                                                                                                                                                                                                                                                               US/08/781,986
                                                                                                                                                                                                                                                                                                                                                                                                                                   3.50 inch, 1.4Mb storage
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                                                                                                                                                                                   PB248PP
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Pred. No. 3.6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    58;
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Query Match

13.4%;

Score 31.2;

DВ

4.

Length 4358;

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Sequence 1, Application US/09918686
Patent NO. 6475739
GENERAL INFORMATION:
APPLICANT: Brunkow, Mary
APPLICANT: Proll, Sean
APPLICANT: Proll, Sean
APPLICANT: Staehling-Hampton, Karen
TITLE OF INVENTION: METHODS FOR IDENTIFY!
TITLE OF INVENTION: GENOMIC DELETIONS
FILE REFERENCE: 240083.515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; LENGTH: 601
; TYPE: DNA
; ORGANISM: Human
US-09-949-016-65898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/949,016
CCURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
VUMBER OF SEQ ID NOS: 207012
SOPTWARE: FRANKSEQ for Windows Version 4.0
SEQ ID NO 65898
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     SUFINE
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                           RESULT 10
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US-09-949-016-65898/c
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Patent No. 6812339

GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                               CURRENT APPLICATION NUMBER: US/09/918,686
CURRENT FILING DATE: 2001-07-30
NUMBER OF SEQ ID NOS: 105
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                                           SOFTWARE: FastSEQ for Windows Version 4.0
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                      62 ТТGAATGAGGCAATCTCTGAGTATGAGACTTCAGAAAAACAATGACTCG 109
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                                                                                                                                                   Karen
DR IDENTIFYING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 30.8; DB 4;
Pred. No. 2;
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Mismatches
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOPTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 16923
LENGTH: 390416
                                                                                                                                                                                                               RESULT 12
US-09-949-016-168377
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Sequence 168377, Application US/09949016

Patent No. 6812339

GENERAL INFORMATION:
APPLICANT: URITER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION PLLE REFERENCE: CLO01307

FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14
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APPLICANT: VENTER, J.
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Best Local Similarity 68.9
Matches 42; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHIENS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
PEATURE:
NAME/KEY: misc_feature
LOCATION: 7043, 8369, 8401
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                            98169
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                                                                                                                                                                                                                                                                                                                                                                                                                                     73 AATCTCTGAGTATGAGACTTCAGAAAACAATGACTCGGGAAGCTACCGCAGATTATTTTA 132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
72; Conserv
                                                                                                                                                                                                                                                                                                                                                   TTTGGCATTGCCTCCATCAGT 153
                                                                                                                                                                                                                                                                                                     TTCCARATTATCTCCCTCAGT 98189
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51.1%; Pred. No. 45
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Pred. No. 23;
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                                                                   AND USES THEREOF
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RESULT 14
US-09-949-016-16483/c
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; ORGANISM: Human
US-09-949-016-12108
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; ORGANISM: Human
US-09-949-016-168377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
        GENERAL INFORMATION:
APPLICANT: VENTER, J.
TITLE OF INVENTION: PO
TITLE OF INVENTION: PO
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 168377
                                                                                                                                                                        Sequence 16483, Application US/09949016 Patent No. 6812339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Patent No. 6812339
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TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CL001307
CHEDENT ASSESSED AND ADDRESS AND ADDRESS
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SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/231,498 PRIOR FILING DATE: 2000-09-08
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mes 50; Conservative
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    Craig et al.
    POLYMORPHISMS IN KNOWN GENES ASSOCIATED
    WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

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Pred. No. 14;
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Pred. No. 3.2;
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OF DETECTION AND USES THEREOF
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US-09-949-016-14856/c
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; ORGANISM: Human
US-09-949-016-16483
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SEQ ID NO 16483
LENGTH: 29628
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                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14856
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Best Local Similarity
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CURRENT ADDITION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
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PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2000-09-08
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                                                                                                                                                                                                                                          NAME/KEY: misc_feature
LOCATION: (1)...(63385)
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                        ORGANISM: Human
                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                     ENGTH: 63385
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                                    36 ATGATAGGGTGGAAGGATTTGAGTTATTGAATGAGGCAATCTCTGAGTATGAGACTTCAG 95
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Pred. No. 31;
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Search completed: May

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Minimum
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*

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US-09-313-294A-6236
US-09-313-294A-5560
US-09-313-294A-5500
US-09-349-016-12501
US-09-949-016-12541
US-09-949-016-13730
US-09-949-016-13770
US-09-949-016-12777
US-09-949-016-12777
US-09-949-016-12777
US-09-949-016-12777
US-09-839-477-7
US-09-839-477-7
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US-09-839-477-7
US-09-839-477-7
US-09-839-477-7
US-09-839-477-7
US-09-949-016-125151
US-09-949-016-125151
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                                 Sequence 6236, Ap
Sequence 4756, Ap
Sequence 13501, A
Sequence 23635, A
Sequence 23635, A
Sequence 13730, A
Sequence 15770, A
Sequence 17777, A
Sequence 7, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 17277, A
Sequence 17391, A
Sequence 17391, A
Sequence 17391, A
Sequence 17391, A
Sequence 1, Appli
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RESULT US-09-3 ; Seque ; Pater ; Pater ; APPI ; APPI ; APPI ; APPI ; TITI ; FILL ; FILL	Qu Ma Bee Oy Oy Oy Oy Db	RESULT US-09- Sequ Fate GENE APP APP APP APP FIL	0 0 0
2 313-294A-4 211 No. 64756, 11 No. 6477 XAL INFORM 11 ALTORMI LI 11 ALTORMI I 11 ALTORMI I 11 ALTORMI SI 12 ALTORMI SI 13 REFERENCI	Query Match Best Local Similarity Matches 99; Conserv (65 ACAAGAATAG B2 ACAAGAATCG 125 TGAAAAGGGC 142 GAGAAAGGG- 185 AGACGAAGCA 185 AGACGAAGCA 191 GCAGCGTGCC 201 GCAGCGTGCC	313-2 ence ence nt No FRAL I LICAN LICAN LICAN LICAN LICAN LICAN LICAN LICAN LICAN LICAN LICAN LICAN LICAN LICAN RENT RENT RENT RENT RENT RENT RENT REN	228 30. 229 30. 31 29.6 13. 32 29.6 12. 33 29.6 12. 34 29.6 12. 36 29.6 12. 37 29.6 12. 39 29.6 12. 40 29.6 12. 41 29.6 12. 42 29.6 12. 43 29.6 12. 44 229.6 12. 45 29.6 12. 46 12. 47 29.6 12. 48 29.6 12. 49 29.6 12. 40 29.6 12. 41 29.6 12. 42 29.6 12. 43 29.6 12. 44 22. 45 29.6 12. 46 12.
756 Application US/09313294A Application US/09313294A ATION: ATION: Co, Laura Y. Co, Laura Y. POLYNUCLECTIDES A S: PL-0017 US	imilarity 66.4%; Pred. No.; Conservative 0; Mismat Acaacaaraagccrigcrigcrigcrigcrif	cation US/(, Raghunathura Y. , Bradley POLYNUCLE -0017 US (1099-05- (1199-05- (1) 7600 (1) 97am	3 15249 4 US-0 1 286 4 US-0 9 1152 5 PCT-1 9 2580 2 US-0 9 2580 4 US-0 9 2580 4 US-0 9 2580 4 US-0 9 2580 4 US-0 9 2589 3 US-0 9 2589 5 PCT-1 9 3532 2 US-0 9 3532 3 US-0 9 3532 3 US-0
.3294A .DES AND POLYPEPTIDES	SCORE 58.6; DB 4; 1 Pred. No. 2.4e-10; 0; Mismatches 49; GATTGGCTGTTAATGGGCAAA;	//09313294A th v. K. LEOTIDES AND POLYPEPTIDES US/09/313,294A 5-14 5-14	US-08-781-986A-102 US-09-313-294A-6591 US-09-222-938A-3 US-09-228-352-3814 US-09-124-001C-592 PCT-US95-05922A-1 US-09-023-655-497 US-09-011-356-7 US-09-011-356-7 US-09-011-356-7 US-09-011-356-7 US-09-012-932-7 US-09-012-932-7 US-09-201-932-7 US-09-201-932-7 US-09-201-932-7 US-09-201-932-7 US-09-201-932-7 US-09-201-932-7 US-09-201-932-7 US-09-201-932-7 US-09-121-931-333 US-09-212-971-7
S DERIVED FROM CORN EAR	25.6%; Score 58.6; DB 4; Length 253; imilarity 66.4%; Pred. No. 2.4e-10; ; Conservative 0; Mismatches 49; Indels 1; Gaps 1 ACAAGAATAGGCCTTGCTGGATTGGCTGTTAATGGCGCAAAAATCTGGCACTCAATATTGCT 124	SS DERIVED FROM CORN EAR	Sequence 102, App Sequence 6591, Ap Sequence 3, Appli Sequence 3814, Ap Sequence 1, Appli Sequence 1, Appli Sequence 7, Appli Sequence 1, Appli

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NAME/KEY: misc_feature

OTHER INFORMATION: Incyte ID No. 6476212 700350270H1

NAME/KEY: unsure

LOCATION: 51

OTHER INFORMATION: a, t, c, g, or other

US-09-313-294A-5508
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LOCATION: 2, 72, 88
OTHER INFORMATION: a,
US-09-313-294A-4756
                                                                                                                                                                                       Matches
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TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FILE REFERENCE: PL-0017 US
CURRENT APPLICATION NUMBER: US/09/313,294A
CURRENT FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 7600
SOFTWARE: PERL Program
SEQ ID NO 5508
LENGTH: 294
TYPE: DNA
ORGANISM: Zea mays
                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local :
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CURRENT FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 7600
SOFTWARE: PERL Program
SEQ ID NO 4756
LENGTH: 277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Lalgudi, Raghunath V. APPLICANT: Ito, Laura Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Zea mays
FEATURE:
NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No. 6476212 700348987H1
                                                                                                                                                                                                   Local Similarity
195 AACCAGGAAGGAAACCTTCAATTTATGGG 223
                                        205 TCCCCATCTCGGTCTACAACAGGA--CAACTCCAAGGTTGATGAGACCGTGCAGCGTGCC
                                                                    135 ТТСССААТТСССОТТААСААСССАТСТССААССТТАТТСССАТАТОВССВАТААССВАССВА
                                                                                                     75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          201 GCAGCGTGCCAAGGCAGAAGGAAACCTTC 229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             125 TGAAAAGGGCTTCCCAATTCCGGTTAACAACGGAACCATTTCCAAGGTTATTGGGCCATA 184
                                                                                                                                                                                     90;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        65 ACAAGAATAGGCCTTGCTGGATTGGCTGTTAATGGGCAAAATCTGGCACTCAATATTGCT 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98;
                                                                                                                                    GCCTTGCTGGATTGGCTGTTAATGGGCAAAATCTGGCACTCAATATTGCTTGAAAAGGGC 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GAGAAAGGG-TTCCCCCATCTCTGTGTACAACAGGACAACCTCCAAGGTGGACGAGACCGT 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGACGAAGCAAACCAGGAAGGAAACCTTC 213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACAAGANTCGGTCTTGCTGGCCTTGCGGTCATGGGGCAGAACCTTGCCCTCAACATTGCA 141
                                                                                                                                                                                  Conservative
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                                                                                                                                                                                              19.1%;
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                                                                                                                                                                         Score 43.8; DB 4;
Pred. No. 3.4e-05;
0; Mismatches 57;
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Pred. No. 5.5e-10;
0; Mismatches 50;
                                                                                                                                                                                                          Length 294;
                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 277;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FROM CORN
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; TYPE: DNA
; ORGANISM: Human
US-09-949-016-12541
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; ORGANISM: Human
US-09-949-016-13501
                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 12541
LENGTH: 133157
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 12541, Application US/09949016
Patent No. 6812339
GEMERAL INFORMATION: 6812139
GEMERAL INFORMATION: GENES ASSOCIATED TITLE OF INVENTION: FOLYMORPHISMS IN KNOWN GENES ASSOCIATED TITLE REFERENCE: CL001307
                                                               Query Match
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                   Matches
                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version SEQ ID NO 13501 LENGTH: 128516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISBASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CLOO1307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR PILING DATE: 2000-09-08
PRIOR RILING DATE: 2000-09-08
PRIOR RILING DATE: 2000-09-08
NUMBER OF SEC. IN SEC. 2000-09-08
NUMBER OF SEC. IN SEC. 2000-09-08
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            Local Similarity 48.9
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nes 87; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16165 ATGCÁÁCAÁGTTGGCAÁTGTAATATTTTÁAAACTGGÁCGÁGÁTGTTGAÁÁÁCATGGTÁ 16222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16105 GTGGAAGGGAATGTGCAAATTATGTCCTGCTATTCTAAGTGAAAGGACTTGCCCAAGGTC 16164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16045 CTTGÁTACCTGÁCCÁAGGTAÁTGGAGAATTGGÁTCCGCCÁGCTCÁTGTGCTGCTTAA 16104
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26 CTTCAAATTTCAGGAGATTAATTATGCGCTCAACCCACAACAACAATAGGCCTTGCTGGA 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      263 AAGGTCGAAGGAAACCTCCCAGTGTTTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGTTAACAACGGAACCATTTCCCAAGGTTATTGGGCCATAAGACGAAGCAAACCAGGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TTGGCTGTTAATGGGCAAAATCTGGCACTCAATATTGCTTGAAAAGGGCCTTCCCAATTCC 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                             14.1%;
48.9%;
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    Score 32.4; DI
Pred. No. 6.5;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Score 32.4; DB; Pred. No. 6.4; 0; Mismatches
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                          DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 4; Length 128516;
    91;
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                                              Length 133157;
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  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                            USES THEREOF
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US-09-513-999C-23635
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                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins Patent No. 6783961
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 23635
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: m=a or c
-09-513-999C-23635
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24 PRIOR APPLICATION NUMBER: US 60/122,487 PRIOR FILING DATE: 1999-02-26 NUMBER OF SEQ ID NOS: 36681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Dumas Milne Edwards, J.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
NAME/KEY: misc_feature
LOCATION: 13
OTHER INFORMATION: w=a c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: m=a or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: h=a or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
LOCATION: 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: 98
OTHER INFORMATION: n=a,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
LOCATION: 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENGTH: 385
                                                                                                                                                                                                                                                                        Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16031 CTTGATACCTGACCAAGGTAATGGAGAATTGGATCCGCCAGCTCATGTGCTGTTGCTTAA 16090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16151 ATGCAACAAGTTGGCAATGTAATATTTTAAAACTGGACGAGATGTTGAAAAACATGGTA 16208
362
                                                                                 302
                                                                                                                                                                   242 ATTTGTTTTTATTGTGCTTGGGGGAGAGGGTATTTTAATATAATTTTTGCCTAAATCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23635, Application US/09513999C
>. 6783961
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                                                                                                                                                                                                                                                                      n 14.0%;
Similarity 51.4%;
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                                                                               GAAGTCCCCTCTGAATGTTAATTTTTAAATGTCAAAATATGATGAACGATATATCTTGAA
                                                                                                                      GAATAGGCCTTGCTGGATTGGCTGTTAATGGGCAAAATCTGGCACTCAATATTGCTTGAA 128
                                                                                                                                                                                                        ATTITITCTGTCATTTGCTTCAAATTTCAGGAGATTAATTATGCGCTCAACCCACAACAA 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTGGCTGTTAATGGGCAAAATCTGGCACTCAATATTGCTTGAAAAGGGGCTTCCCAATTCC 145
AGTGAGATTGCAATATGCTTAAAC 385
                                     AAGGGCTTCCCAATTCCGGTTAAC 152
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                                                                                                                                                                                                                                                Score 32; DB 4; Length 385; Pred. No. 0.48; 0; Mismatches 70; Indels
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                                                                          ; FEATURE:
, NAME/KEY: misc_feature
; LOCATION: (1)...(163181)
; OTHER INFORMATION: n = A,T,C
US-09-949-016-13730
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PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
Query Match
Best Local S
Matches 66
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 79076
                                                                                                                                                                                                                                  SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 13730
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Patent No. 681233
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Patent No. 6812339
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                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR PILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/949,016
                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Human
                                                                                                                                                                          ORGANISM: Human
                                                                                                                                                                                                 LENGTH: 163181
TYPE: DNA
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1 Similarity
66; Conserv
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                    13.9%;
53.7%;
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Pred. No. 0.71;
0; Mismatches 27;
                    Score 31.8;
Pred. No. 12;
  Mismatches
                                         DB
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  57;
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OF DETECTION AND USES THEREOF
                                     Length 163181;
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103 AAATCTGGCACTCAATATTGCTTGAAAAGGGCTTCCCAATTCCGGTTAACAACGGAACCA 162

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US-09-949-016-14033/c

Sequence 14033, Application US/09949016

Patent No. 6812339
                              CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR PRIOR PRICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
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SEQ ID NO 15770
LENGTH: 211049
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Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISBASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                          APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
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Best Local
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
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NUMBER OF SEQ ID NOS: 207012
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                                                                                                                                                                                                                                                                                                                                                                                                                            22794 ATTTCAAAATGTATTCACATTTATGGGTAAAAACATTACCACT 22752
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22914 AAATGTTGTATTAATAGAGGCTTTGTACAAATTTGAGAGAAATGAGGCACTGCCTAGTGC 22855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 122 GCTTGAAAAGGGCTTCCCAATTCCGGTTAACAACGGAACCATT 164
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ch 13.9%;
1 Similarity 49.7%;
81; Conservative (
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                    FastSEQ for Windows Version 4.0
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Pred. No. 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches 82;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 4;
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Sequence 7, Application US/09839477
PATENT NO. 6723895
GENERAL INFORMATION:
APPLICANT: Shorrosh, Basil S.
APPLICANT: DeBonte, Lorin R.
TITLE OF INVENTION: PLANTS CONTAIN
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Best Local !
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; NAME/KEY: misc_feature
; LOCATION: (1)...(828152)
; OTHER INFORMATION: n = A
US-09-949-016-12777
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; NAME/KEY: misc_feature
; LOCATION: (1)...(784019)
; OTHER INFORMATION: n = A,T,C or G
US-09-949-016-14033
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                                                                                                                                                                                                                                                                                                                                             APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR TELLING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                         Matches 48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 12777,
                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Human
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 14033
LENGTH: 784019
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Best Local Similarity
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   470711
                                                              470771 AAAGCAGAATITTTCTTAGATTTGCTTCATATTGGAAGTAAATATGTGTGGGGCTCAATAT 470712
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                                62 ACAACAAGAATAGGC 76
                                                                                  48;
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                                                                                                                                                         Similarity
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6812339
ACAACTGAAATTAGC 470697
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                                                                                                                                       Conservative
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                                                                                                                                                      Score 31.8;
Pred. No. 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 31.8; DE Pred. No. 25; 0; Mismatches
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                                                                                                                                     Mismatches
                                                                                                                                                                   DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 4;
                                                                                                                                     27;
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                                                                                                                                                                    Length 828152;
                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 784019;
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                                                                                                                                   0,
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                                                                                                                                Gaps
                                                                                                                                  <u>.</u>
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S.
R.
CONTAINING A CYTOSOLIC ACETYL

TITLE OF INVENTION: COA-CARBOXYLASE FILE REFERENCE: 07148-094001 CURRENT APPLICATION NUMBER: US/09/839,477 CURRENT FILING DATE: 2001-04-20

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                                                                                                                               RESULT 14
US-09-949-016-16480
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US-09-614-221A-346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-614-221A-346
      GENERAL INFORMATION:
APPLICANT: VENTER, J.
TITLE OF INVENTION: PO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 346, Application US/09614221A Patent No. 6723837
                                                                                      Sequence 16480, Application US/09949016 Patent No. 6812339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 346
LENGTH: 2523
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 7
LENGTH: 7151
                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/198,794
PRIOR FILING DATE: 2000-04-20
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Yu, Jaehyuk
APPLICANT: Kishore, Ganesh M.
APPLICANT: Kishore, Ganesh M.
TITLE OF INVENTION: WICLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED
TITLE OF INVENTION: WITH STEROL SYNTHESIS AND METABOLISM
FILE REFERENCE: 16516.075
CURRENT APPLICATION NUMBER: US/09/614,221A
CURRENT FILING DATE: 2000-07-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/142,981
PRIOR FILING DATE: 1999-07-12
NUMBER OF SEQ ID NOS: 626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Karunanandaa, Balasulojini
APPLICANT: Yu, Jaehyuk
APPLICANT: Kishore, Ganesh M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Medicago sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: CDS
LOCATION: (1)...(6771)
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                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5807 CTGATTCTGCGACCAAGACGGCCCAAGCGATATTGGATTTCAACAGAGAAGAACTCCCAC 5866
                                                                                                                                                                                                                   2365
                                                                                                                                                                                                                                                                                                 2305 GCATGGGACTTGCTGGATTTAAAATTAGTACTGCGAAATGTACCTATATTACCAATTTTG 2364
                                                                                                                                                                                                                                                        129 AAGGGCTTCCCAATTCCGGTTAACAACGGAACCATTTCCAAGGTTATT 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            114 TCAATATTGCTTGAAAAGGGCTTCCCAATTCCGGTTAACAACGGAACCATTTCCAAGG 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       64; Conservative
                                                                                                                                                                                                                                                                                                                                          69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              54 CTCAACCCACAACAAGAATAGGCCTTGCTGGATTGGCTGTTAATGGGCAAAATCTGGCAC 113
                                                                                                                                                                                                                                                                                                                                                                                   60;
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                                                                                                                                                                                                                                                                                                                                        GAATAGGCCTTGCTGGATTGGCTGTTAATGGGCAAAAATCTGGCACTCAATATTGCTTGAA 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TTTTCATTATCGCAAACTGGAGAGGCTTTTCAGGTGGACAAAGGGACCTTTTTGAAGG
                                                                                                                                                                                                                 AACGGTCAGCCAÁTTCATGGAÁACAAGGTÁAGAATCAACAAGGTTÁTT
                                                                                                                                                                                                                                                                                                                                                                                   Conservative

    Craig et al.
POLYMORPHISMS IN KNOWN GENES ASSOCIATED
WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

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                                                                                                                                                                                                                                                                                                                                                                                                                       DB 4;
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Sequence 1, Application.

Patent No. 6506581

Patent No. 6506581

APPLICANT: Fleischmann et al.

APPLICANT: Fleischmann et al.

TITLE OF INVENTION: The Nucleotide sequence of the Haemophilus influenzae Rd

The Haemophilus influenzae Rd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 16480
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE: misc_feature
NAME/KEY: misc_feature
LOCATION: (1)...(225127)
OTHER INFORMATION: n = A,T,C or G
                       INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENGTH:
                                                                                                   PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/476,102
EILING DATE: JUN-5-1995
ATTORNEY/AGENT INFORMATION:

NAME: Michelle S. Marke
REGISTRATION NUMBER: 41,971
REFERENCE/DOCKET NUMBER: PB186P3
TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
nes 63; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16067 TTÄÄAAGACÁCÁGAGTGGCAAGCTÁGÄTAAAAAGACAAGACCCAACCATTTGCTGTCTTC 16126
                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: ASCII Text CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3 1/2 inch diskette
COMPUTER: Dell Pentium
OPERATING SYSTEM: MS DOS v6.22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93 TTAATGGGCAAAATCTGGCACTCAATATTGCTTGAAAAGGGCTTCCCAATTCCGGTTAAC 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               225127
                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/557,884
FILING DATE: 25-Apr-2000
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Rockville
STATE: MD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATGAGACCAATTTCACATGTAATGGTACCCACAGGCGCAAAGTAAAAGGATGGAGA 16182
                                                                TELEFAX: 301-309-8439
                                                                                     TELEPHONE: 301-309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 9410 Key West Avenue
LENGTH: 1830121 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20850
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Pred. No. 22;
0; Mismatches
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Title:
Perfect score:
Sequence:
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Maximum DB seq length: 2000000000
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OM nucleic -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Database :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        May 4, 2005, 12:06:42; Search time 259.813 Seconds (without alignments) 5365.716 Million cell updates/sec
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Gapop 10.0 , Gapext 1.0
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229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Published_Applications_NA:*
| (ggn2_6/ptcdata/1/pubpna/US07_N: /ggn2_6/ptcdata/1/pubpna/US06_
| (ggn2_6/ptcdata/1/pubpna/US06_)
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Copyright (c) 1993 - 2005 Compugen Ltd.
'cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
(cgn2_6/ptodata/1/pubpna/US11_NEW_PUB.seq:*
(cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
(cgn2_6/ptodata/1/pubpna/US60_PUBCOMB.seq:*
                                                                                                      jn2_6/ptodata/1/pubpna/US09C_PUBCOMB.seq:*
jn2_6/ptodata/1/pubpna/US09 NEW PUB.seq:*
jn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
jn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
jn2_6/ptodata/1/pubpna/US10C_PUBCOMB.seq:*
jn2_6/ptodata/1/pubpna/US10D_PUBCOMB.seq:*
jn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
jn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
jn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
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1/pubpna/PCT_NEW_PUB.seq:*
1/pubpna/US06_NEW_PUB.seq:*
1/pubpna/US06_PUBCOMB.seq:*
1/pubpna/US07_NEW_PUB.seq:*
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/pubpna/USO8_NEW_PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Result No.

esult No.	Score	Query Match	Query Match Length DB	DB	ID	Description	
ם	137.8	60.2	4825	17	US-10-424-599-59918	Sequence 59918, A	
2	136.2	59.5	2691	17	US-10-424-599-59924		
ω	126.6	55.3	578	17	US-10-424-599-59920		
4.	117	51.1	1737	17	US-10-425-114-12849		
տ	81	35.4	1627	17	US-10-425-114-10641	Sequence 10641, A	
6	18	35.4	1838	17	US-10-424-599-76904	Sequence 76904, A	
7	74.2	32.4	1785	18	US-10-739-930-1129	Sequence 1129, Ap	
æ	73.8	32.2	375	17	US-10-424-599-98726		
9	63.4	27.7	1125	18	US-10-425-115-44920	Sequence 44920, A	
10	63.4	27.7	1953	18	US-10-767-701-14931	Sequence 14931, A	
11	62.8	27.4	234	11	US-09-732-627A-2087	Sequence 2087, Ap	

ALIGNMENTS

120 2449	CAATAT	GCACTO	AAATCTG	ATGGGC	GCTGTT!	GCTGGATT	CACAACAAGAATAGGCCTTGCTGGATTGGCTGTTAATGGGCAAAATCTGGCACTCAATAT	61 CACAACAAGAATAGGCCTTGCTGGATTGGCTGTTAATGGGCAAAATCTGGCACTCAATAT		음 성
60 2389		ATGCGCT	ATTAATTA \TCAATTA	CAGGAGI	CAAATTI	CATTIGCT	CAGACCTTATTTTTTCTGTCATTTGCTTCAAATTTCAGGAGATTAATTA		y 1 0 2331	유 성
2;	Gaps		Length 4825; Indels 2;	7;	ore 137.8; [ed. No. 9.9e- Mismatches	Score 137.8; DB 1 Pred. No. 9.9e-34; 0; Mismatches 42	60.2%; 80.8%; vative	Query Match Best Local Similarity 80. Matches 185; Conservative	Query Match Best Local Matches 18	
				.7C.1		PAT_MRT3	: 285684 lax Clone ID:	NUMBER OF SEQ ID NOS: 285684 EQ ID NO 59918 LENGTH: 4825 TYPE: DNA ORGANISM: Glycine max PERATURE: OTHER INFORMATION: Clone ID: PAT_MRT3847_25117C.1 10-424-599-59918	NUMBER OF SEQ ID SEQ ID NO 59918 LENGTH: 4825 TYPE: DNA ORGANISM: Glyci ORGANISM: INFORMATI US-10-424-599-59918	g
iated With	s Associated t	lecules	Acid Molecules and Other Molecules Uses Thereof for Plant Improvement 10/424,599	and Ot	olecules nereof f	.c Acid Mo nd Uses T) 3/10/424,	David K ua wei Soy Nucleic Plants and 1 (53223)B NUMBER: US/1 2003-04-28	APPLICANT: KOVALIC DAVIG K APPLICANT: Zhou Yihua APPLICANT: Zhou Yihua APPLICANT: Cao Yongwei TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53223)B CURRENT APPLICATION NUMBER: US/10/424,599 CURRENT FILING DATE: 2003-04-28	APPLICANT: APPLICANT: APPLICANT: TITLE OF I TITLE OF I TITLE REFER CURRENT AP	
						10424599	cation US/ 40031072A1	SULT 1 10-424-599-59918 Sequence 59918, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION: APPLICANT: LA ROSA Thomas J APPLICANT: LA ROSA Thomas J	RESULT 1 US-10-424-599-59918 US-10-424-599-59918, A ; Sequence 59918, A ; Publication No. U ; GENERAL INFORMATI ; APPLICANT: La Ro ; APPLICANT: La Ro	· · · · · · · · · · · · · · · · · · ·

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RESULT 3
US-10-424-599-59920
; Sequence 59920, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
   APPLICANT: La Rosa Thomas J
   APPLICANT: Cao Yongwei
   TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
   TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
   FILE REFERENCE: 38-21(5323)B
   CURRENT APPLICATION UNUMBER: US/10/424,599
   NUMBER OF SEQ ID NOS: 285684
   SEQ ID NO 59920
   LENGTH: 578
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APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)B
FULE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 59924
LENGTH: 2691
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Best Local
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                 ORGANISM: Glycine max
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TĠĊ-TĠAGĀĀĀĠĠĊTŤTĊĊĊĀŤŤTĊTĠŤŤTĀTĀĀĊCĠĀĀĊĊĀCŤŤĊĊĀĀĠĠŤŤGĀŤĠĀĠĀ 2508
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US-10-425-114-12849
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Best Local Similarity
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APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, Davi
APPLICANT: Screen, Steve
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Publication No. US20040034888A1
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Best Local Similarity
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NAME/KEY: unsure
LOCATION: (1)...(578)
OTHER INFORMATION: unsure at all n locations
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                             CAATTTATGGGGAACAA 229
CCAGTTTATGGCTÁCCÁ 237
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APPLICANT: KOWAIIC, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5313)B
CURRENT TAPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 12849
LENGTH: 1737
TYPE: DNA
ORGANISM: Glycine max
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                                      394 CTGTAGAACGAGCAAAAACAAGAAGGAAATCTTCCAGTTTATGGCTACCA
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                                                                                    TTATGGGCCAAAATCTGGCACTCAATATTGC-TGAGAAAGGCTTTCCCCATTTCTGTTTAC
                                                                                                                      TTAATGGGCAAAATCTGGCACTCAATATTGCTTGAAAAGGGCTTCCCAATTCCGGTTAAC 152
                                                                                                                                                                              TATCTTGAGATTAATTATG-GCTCAACCCACAACAAGAATAGGCCTTGCTGGATTGGCTG
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                                                                                                                                                     ; OTHER INFORMATION: Clone ID: PAT_MRT3847_40459C.1
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CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 10641
LENGTH: 1627
TYPE: DNA
ORGANISM: Glycine max
                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 76904
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
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ORGANISM: Glycine max
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                                                                                        Local
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Similarity 72.1%;
                                                                                      Similarity
ACAAGAATAGGCCTTGCTGGATTGGCTGTTAATGGGCAAAATCTGGCACTCAATATTGCT 124
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Kovalic, David K.
Screen, Steven E
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                                                                  Conservative
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O; Mismatches
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Pred. No. 1.7e-15;
0; Mismatches 45;
                                                                                   DB 17; Length 1838;
1.8e-15;
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; OTHER INFORMATION: Clone ID: PAT_MRT3847_60162C.1 US-10-424-599-98726
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                                                                                                                                                                                    APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)B
FULE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 98726
LENGTH: 375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 98726, Application US/10424599 Publication No. US20040031072A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 117; Conserv
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SEQ ID NO 1129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 38-21(53377)B
CURRENT APPLICATION NUMBER: US/10/739,930
CURRENT FILING DATE: 2003-12-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: KOVALIC, DAVID K.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH
TITLE OF INVENTION: PLANTS AND USES THEREOF FOR PLANT IMPROVEMENT
                                                                                                                       ORGANISM: Glycine max FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE: OTHER INFORMATION: Clone ID: ARATH-23APR03-CLUSTER6852_1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENGTH: 1785
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   113 CTCAATATTGCTTGAAAAGGGCTTCCCAATTCCGGTTAACAACGGAACCATTTCCAAGGT 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     224 ACAAGAATAGGCCTTGCTGGCCTGGCTGTGATGGGCCAAAAACCTTGCTCTCAACATTGC- 282
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGACGAAGCAAACCAGGAAGGAAACCTTCAATTTATGGGGAACAA 229
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TATTGGGCCATAAGACGAAGCAAACCAGGAAGGAAACCTTCAATTTATGGG 223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTCAACATTGCA-GAGAAAGGCTTCCCAATCTCTGTTTACAACAGAACTACTTCCAAAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GCTGTTCAACCTACAAGAATAGGTCTAGCTGGACTTGCTGTGATGGGTCAGAACCTAGCT 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TGAAAGAGCAAAACAAGAAGGAAACCTTCCTGTGTATGGCTACCA 387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGAAAAGGGCTTCCCAATTCCGGTTAACAACGGAACCATTTCCAAGGTTATTGGGCCATA 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGAGAAAGGCTTTCCCATTTCTGTCTACAACCGGACCGCGTCCAAGGTTGATGAGACAGT 342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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  32.2%;
63.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 74.2; DB 18;
Pred. No. 2.9e-13;
  Score 73.8; DB 17;
Pred. No. 1.9e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                         Length 375;
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US-10-767-701-14931
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; Sequence 44920, Application US/10425115

; Publication No. US20040214272A1
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                                                                                                        Sequence 14931, Application US/10767701 Publication No. US20040172684A1 GENERAL INFORMATION:
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APPLICANT: Kovalic, David K.

APPLICANT: Zhou, Yihua

APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants

FILE REFERENCE: 38-21 (53222)B

CURRENT APPLICATION NUMBER: US/10/425,115

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 369326
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof For Plant Improvement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 44920
LENGTH: 1125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Zea mays
FEATURE:
NAME/KEY: unsure
LOCATION: (1) . (1125)
OTHER INFORMATION: unsure at all n locations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                       185 AGACGAAGCAAACCAGGAAAGGAAACCTTC 213
                                                                                                                                                                                                                                                                                                                                                                                      166 ACCAGAATTGGTCTTGCTGGCCTCGCCGTCATGGGGCAGAACCTTGCCCTCAACATTGCA 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           102;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                              TGAAAAGGGCTTCCCCAATTCCGGTTAACAACGGAACCATTTCCAAGGTTATTGGGCCATA 184
                                                                                                                                                                                                                                           TCAGCGCGCCAAGCTAGAAGGAAACCTTC
                                                                                                                                                                                                                                                                                                             -GAGAAAGGCTTCCCCATCTCTGTCTACAACAGGACCACCTCCAAGGTCGATGAGACTGT 284
                                                                                                                                                                                                                                                                                                                                                                                                                          ACAAGAATAGGCCTTGCTGGATTGGCTGTTAATGGGCAAAATCTGGCACTCAATATTGCT 124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGCTTGAAAAGGGCTTCCCAAGTCCGGTTAACAACGGAACCATTTCCCAAGGTTATTGGGC 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CGCATCATGAATTCCACTTGCTGGGTAGCTTCTTATGGGCCAAACCCCTGGCGCTCACTAT 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TGATCCGAACGGCTGTCCCATTAATGATAACAACTCTAACCACTTCCAATGCTGCTGAGA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CACATCTTCTCTGTTGTGATTTGCTACCCTTTTTAGGAGAATGATTAT-TGCTCCTCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 63.4; DB 18;
Pred. No. 7.7e-10;
0; Mismatches 46;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 1125;
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OTHER INFORMATION: Clone ID: LIB3493-030-P1-M1-G8
US-09-732-627A-2087
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US-09-732-627A-2087
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                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/732,627A
CURRENT FILING DATE: 2000-12-08
NUMBER OF SEQ ID NOS: 4930
SEQ ID NO 2087
LENGTH: 234
                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                              Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2087, Application US/09732627A
Publication No. US20040123338A1
GENERAL INFORMATION:
APPLICANT: Fincher, Karen L.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with
TITLE OF INVENTION: plants
FILE REFERENCE: 38-21(51770) B
FILE REFERENCE: 38-21(51770) B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 14931
LENGTH: 1953
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 38-21(53535)B
CURRENT APPLICATION NUMBER: US/10/767,701
CURRENT FILING DATE: 2004-01-29
NUMBER OF SEQ ID NOS: 63128
                                                                                                                                                                                                                                                                                                                                             NAME/KEY: unsure
                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Gossypium
                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                           LOCATION: (1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Sorghum bicolor FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
     197
                                  179
                                                                     137
                                                                                            119 ATTGCTTGAAAAGGGCTTCCCCAATTCCCGGTTAACAACGGAACCATTTCCCAAGGTTATTGG 178
                                                                                                                                                   59 CCCACAACAAGAATAGGCCTTGCTGGATTGGCTGTTAATGGGCAAAATCTGGCACTCAAT 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              185 AGACGAAGCAAACCAGGAAGGAAACCTTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          332 GCAGCGTGCCAAGGCAAAGGAAACCTTC
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                                                                                                                                                                                                                            Similarity
                              GCCATAAGACGAAGCAAACCAGGAAGGAAACCTT 212
                                                          TTTTTTGĀTĀĀĀĠĠĊTTĊĊĊĊĀĀTĀĊTĠTTTĀĊĀĀĊĀĠĀĀCTĀCCTČĊĀĀĀĠTCGĀTĠĀ
GACTGTTGÁAAGAGCTÁAACAGGÁAGGGGGTCTT
                                                                                                                             CCACCTACAAGAATAGGACTIGCTGGTTTGGCTGTTATGGGCCGGGGGGGCTTGCGCTTAA 136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACAAGAATTGGTCTTGCTGGCCTTGCGGTCATGGGGCAGAACCTTGCCCTCAACATTGCA 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -GAGAAAGGCTTCCCCATCTCTGTCTACAACAGGACAACCTCCAAGGTTGATGAGACCGT 331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TGAAAAGGGCTTCCCAATTCCGGTTAACGAACCATTTCCAAGGTTATTGGGCCATA 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACAAGAATAGGCCTTGCTGGATTGGCTGTTAATGGGCAAAATCTGGCACTCAATATTGCT 124
                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                (234)
                                                                                                                                                                                                                      27.4%;
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68.5%;
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                                                                                                                                                                                                                     Score 62.8; DB 11
Pred. No. 5.7e-10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 63.4; DB 18
Pred. No. 9.9e-10;
                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          360
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                                                                                                                                                                                                                                     DB 11;
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                                                                                                                                                                                                      57;
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                                                                                                                                                                                                                                     Length 234;
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US-10-424-599-60294

Sequence 60294, Publication No.

Application US/10424599 US20040031072A1

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US-10-424-599-60294
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TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 60294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 98959, Application US/10437963 Fublication No. US20040123343A1
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                                                                                                                                                                    Query Match
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SEQ ID NO 98959
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: La Rosa, Thomas J. APPLICANT: Kovalic, David K.
                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53221)B
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APPLICANT: Kovalic David
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                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                       ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                   TYPE: DNA
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125 TGAAAAGGGCTTCCCAATTCCGGTTAACAACGGAACCATTTCCAAGGTTATTGGGCCATA 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          125 TGAAAAGGGCTTCCCAATTCCGGTTAACAACGGAACCATTTCCAAGGTTATTGGGCCATA 184
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                                                                                                                                  106; Conservative
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                                                                                                                                                                                                                                                                                                   1831
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Li, Ping
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Barbazuk, Brad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wu, Wei
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                                                                                                                                0; Mismatches
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Pred. No.
                                                                                                                                                 Score 60.2; DB 18; Pred. No. 1.1e-08;
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                                        ; OTHER INFORMATION: Clone ID: MRT4577_183003C.1 US-10-425-115-91011
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US-10-425-115-91011
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US-10-425-114-12436
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                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 91011
LENGTH: 637
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CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 12436
                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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Query Match
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
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                                                                                ORGANISM: Zea mays
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 283 GCAGCGTGCCAAGGCAGAAGGAAACCTTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        224 GAGAAAGGG-TTCCCCATCTCTGTGTACAACAGGACAACCTCCAAGGTGGACGAGACCGT 282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            125 TGAAAAGGGCTTCCCAATTCCGGTTAACAACGGAACCATTTCCAAGGTTATTGGGCCATA 184
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99; Conser
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Screen, Steven E
Tabaska, Jack E
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Pred. No. 2.1e-08;
0; Mismatches 49;
Score 58.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 311
DB 18;
Length 637;
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Perfect score:
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Gapop 10.0 , Gapext 1.0
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229
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1: /cgn2_6/ptodata/1/pna/
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/ Cgn2_6/ptodata/1/pna/US081_COMB.seq:*
/ Cgn2_6/ptodata/1/pna/US081_COMB.seq:*
/ Cgn2_6/ptodata/1/pna/US082_COMB.seq:*
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/ Cgn2_6/ptodata/1/pna/US084_COMB.seq:*
/ Cgn2_6/ptodata/1/pna/US085_COMB.seq:*
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/ Cgn2_6/ptodata/1/pna/US086_COMB.seq:*
/ Cgn2_6/ptodata/1/pna/US086_COMB.seq:*
/ Cgn2_6/ptodata/1/pna/US089_COMB.seq:*
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474 (-cgm2-6/ptodata/
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Score
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Match
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Result No.

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US-09-333-535A-6002
US-09-371-146A-181981
US-09-684-617-43713
US-09-684-617-43713
US-09-684-617-43713
US-09-985-678-181981
US-09-985-678-19274
US-09-987-678-109274
US-09-987-4708A-18187
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US-09-987-703-14731
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US-09-985-678-235976
US-09-985-678-235976
US-09-09-615-606A-10170
US-09-09-615-606A-10170 Sequence e 181981, Appl e 26002, Appl e 181981, Appl e 2619274, Appl e 109274, e 109274, e 109274, e 116027, Appl e 1160

ALIGNMENTS

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APPLICANT: Chelki, Notated Plant Genes
APPLICANT: Liu, Jingdong Annotated Plant Genes
FILE REFERENCE: 38-21(1597)B
CURRENT APPLICATION NUMBER: US/09/304,517A
CURRENT FILING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 295529
SEQ ID NO 191981
LENGTH: 229
TYPE: DNA
ORGANISM: Glycine max
US-09-304-517A-181981
100.0%; Score 229; DB 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 i NUMBER OF SEQ ID NOS
SEQ ID NO 27
LENGTH: 229
TYPE: DNA
ORGANISM: Glycine π
US-09-300-482-27
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; GENERAL INFORMATION:
; APPLICANT: Cheikh, Nordine
; APPLICANT: Liu, Jingdong
; APPLICANT: Liu, Jingdong
; TITLE OF INVENTION: Nucleic Acid Molecules And Ot;
; TITLE OF INVENTION: Phosphogluconate Pathway
; FILE REFERENCE: 04983.0031.US01/38-21(15365)B
; CURRENT APPLICATION UMBER: US/09/300,482
; CURRENT FILING DATE: 1999-04-28
; NUMBER OF SEQ ID NOS: 699
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; Sequence 181981, Application US/09304517A
; GENERAL INFORMATION:
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US-09-300-482-27
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Best Local S
Matches 229
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                                                    CACAACAAGAATAGGCCTTGCTGGATTGGCTGTTAATGGGCAAAATCTGGCACTCAATAT
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                                                                                                                                                                      Conservative
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                                                                                                                                                                100.0%; Score 229; DB 20;
100.0%; Pred. No. 1.3e-60;
cive 0; Mismatches 0;
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Pred. No. 1.3e-60;
; Mismatches 0;
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; APPLICANT: Liu, Jingdong
; TITLE OF INVENTION: ANNOTATED PLANT GENES
; FILE REFERENCE: 38-21(15097)C
; CURRENT APPLICATION NUMBER: US/09/371,146A
; CURRENT FILING DATE: 1999-08-09
; PRIOR APPLICATION UNBER: US 09/304,517
; PRIOR FILING DATE: 1999-05-06
; PRIOR FILING DATE: 1999-05-06
; MUMBER OF SEQ ID NOS: 294310
; SEQ ID NO 101981
; LENGTH: 229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; LENGTH: 229
; TYPE: DNA
; ORGANISM: Glycine max
; OTHER INFORMATION: Clc
US-09-333-535A-6002
                                                                                                                                                               ; TYPE: DNA; ORGANISM: Glycine max US-09-371-146A-181981
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                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 181981, Application US/09371146A GENERAL INFORMATION:
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Best Local Similarity
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                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/333,535A CURRENT FILING DATE: 1999-06-14 NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: La Rosa, Thomas J.
APPLICANT: Thompson, Michael D.
TITLE OF INVENTION: Nuclei
TITLE OF INVENTION: Plants
78-21(1540
                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Cheikh, Nordine APPLICANT: Liu, Jingdong
                                                                                                             Local Similarity
                                                                                            229;
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                                        CATAAGACGAAGCAAACCAGGAAGGAAACCTTCAATTTATGGGGAACAA 229
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                                                                                            Conservative
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                                                                                                         100.0%;
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                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                       Score 229; DB 21;
Pred. No. 1.3e-60;
); Mismatches 0;
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Pred. No. 1.3e-60;
; Mismatches 0;
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                                                                                          Indels
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                                   ; TYPE: DNA; Glycine max; ORGANISM: Glycine max US-09-654-617-43713
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                                                                                   FILE REFERENCE: 38-21(15097)D
CURRENT APPLICATION NUMBER: US/09/654,617
CURRENT FILING DATE: 2000-09-05
NUMBER OF SEQ ID NOS: 463173
SEQ ID NO 43713
LENGTH: 229
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CURRENT APPLICATION NUMBER: US/09/552,086
CURRENT FILING DATE: 2000-04-19
NUMBER OF SEQ ID NOS: 27278
SEQ ID NO 26409
LENGTH: 229
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Matches 229; Conserv
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 Query Match
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APPLICANT: Liu, Jir
TITLE OF INVENTION:
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APPLICANT: Byrum, Joseph R.
APPLICANT: Coombs, Brian E.
APPLICANT: La Rosa, Thomas J.
TITLE OF INVENTION: Nucleic A
TITLE OF INVENTION: Plants
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ORGANISM: Glycine max
OTHER INFORMATION: Clone
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 100.0%;
                                                                                                                                                                                             Annotated Plant Genes
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; Pred. No. 1.3e-60;
0; Mismatches 0;
 Score 229;
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APPLICANT: Liu, Jingdong
APPLICANT: Liu, Jingdong
TITLE OF INVENTION: AND Annotated Plant Ger
FILE REFERENCE: 38-21(1597)D
CURRENT APPLICATION NUMBER: US/09/684,016
CURRENT FILING DATE: 2000-10-10
PRIOR APPLICATION NUMBER: US 09/654,61
PRIOR APPLICATION NUMBER: 2000-09-05
NUMBER OF SEQ ID NOS: 463173
LENGTH: 229
TYPE: DNA
ORGANISM: Glycine max
US-09-684-016-43713
                         Sequence 181981, Application US/09985678
GENERAL INFORMATION:
APPLICANT: Chekk, Nordine
APPLICANT: Chekk, Nordine
APPLICANT: Liu, Jingdong
TITLE OF INVENTION: Annotated Plant Genes
FILE REFERENCE: 16517.255/38-21(15097) F
CURRENT APPLICATION NUMBER: US/09/985,678
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: US 09/304,517
PRIOR FILING DATE: 1999-05-06
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US-09-684-016-43713
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Best Local :
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              SEQ ID NOS: 295529
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; Mismatches 0;
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; ORGANISM: Glycine
US-09-985-678-181981
                                                                                                                               TELEPAX: (314) 737-604/
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 305 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 829, Application US/60089810
GENERAL INFORMATION:
APPLICANT: Joseph R. Byrum
APPLICANT: Michael D. Thompson
TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules
TITLE OF INVENTION: Associated With Plants.
NUMBER OF SEQUENCES: 5409
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ORIGINAL SOURCE:
ORGANISM: Glycine max
DEVELOPMENTAL STAGE: F
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NAME: Lavin Jr., Lawrence M.
REGISTRATION UNMBER: 30,768
REFERENCE/DOCKET NUMBER: 38-21(15395)A
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette (3.5in, 1.4
COMPUTER: ISM PC/XT/AT, ISM PS/2
OPERATING SYSTEM: Windows 95/NT
SOFTWARE: WinZip 6.3
CURRENT APPLICATION DATA:
                                                                       MOLECULE TYPE:
HYPOTHETICAL:
                                                         ANTI-SENSE:
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CORRESPONDENCE ADDRESS:
                                                                                                                       STRANDEDNESS:
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STREET: BB4F
STREET: 700 Chesterfield Parkway North
CITY: St. Louis
                                                                                                                                                                                                                TELEPHONE: (314) 737-6670 TELEFAX: (314) 737-6047
                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
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BB4F
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Pred. No. 1.3e-60;
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SEQ ID NO 70

LENGTH: 261

TYPE: DNA

ORGANISM: Glycine max
US-09-300-482-70
                                                                 RESULT 11
US-09-304-517A-109274
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                ; Sequence 109274, Application US/09304517A; GENERAL INFORMATION:
; APPLICANT: Cheikh, Nordine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 70, Application US/09300482
GENERAL INFORMATION:
APPLICANT: Cheikh, Nordine
APPLICANT: Liu, Jingdong
APPLICANT: Liu, Jingdong
TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
TITLE OF INVENTION: Phosphogluconate Pathway
FILE REFERENCE: 04983.0031.US01/38-21(15365)B
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Best Local S
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CURRENT FILING DATE: 1999-04-28
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LIBRARY: SOYMON028
CLONE: 701097624H1
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Local Similarity 83.7%;
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                                                                                                                                                                                                                                                                                 CACAACAAGAATAGGCCTTGCTGGATTGGCTGTTATGGGCCAAAATCTGGCACTCAATAT 125
                                                                                                                                                                                                                                                                                                                  CACAACAAGAATAGGCCTTGCTGGATTGGCTGTTAATGGGCAAAATCTGGCACTCAATAT 120
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                                                                                                                                        CAGTAGAACGAGCAAAACAAGAAGGAAATCTTCCAGTTTATGGCTACCA 233
                                                                                                                                                                                                              TGC-TGAGAAAGGCTTTCCCATTTCTGTTTACAACCGAACCACTTCCAAGGTTGATGAGA
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Liu, Jingdong
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Pred. No. 4.5e-37;
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; SEQ ID NO 109274
; LENGTH: 261
; TYPE: DNA
; ORGANISM: Glycine max
US-09-304-517A-109274
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APPLICANT: Liu, Jingdong
TITLE OF INVENTION: ANNOTATED PLANT GENES
FILE REFERENCE: 38-21(15097)C
CURRENT APPLICATION NUMBER: US/09/371,146A
CURRENT FILING DATE: 1999-08-09
FRIOR APPLICATION NUMBER: US 09/304,517
PRIOR APPLICATION NUMBER: US 09/304,517
PRIOR FILING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 294310
SEQ ID NO 109274
LENGTH: 261
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US-09-371-146A-109274
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Best Local (
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185
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                                                                                                                              66 CACAACAAGAATAGGCCTTGCTGGATTGGCTGTTATGGGCCAAAATCTGGCACTCAATAT 125
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                              CATAAGACGAAGCAAACCAGGAAGGAAACCTTCAATTTATGGGGAACAA
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85.2%;
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                                                                                                                                                                                                                                                             Score 153.8; DB 2
Pred. No. 4.5e-37;
0; Mismatches 32
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Pred. No. 4.5e-37
0; Mismatches 3:
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RESULT 13

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APPLICANT: Cheikh, Nordine
APPLICANT: Liu, Jingdong
TITLE OF INVENTION: Annotated Plant Genes
FILE REFERENCE: 16517.355/38-21(1597) F
CURRENT APPLICATION NUMBER: US/09/985,678
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: US 09/304,517
PRIOR FILING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 295529
SEQ ID NO 109274
LENGTH: 261
TYPE: DNA
ORGANISM: Glycine max
US-09-985-678-109274
                                                                                                                                                                                                                                                                                                           APPLICANT: Byrum, Joseph R.
APPLICANT: La Rosa, Thomas J.
APPLICANT: La Rosa, Thomas J.
APPLICANT: Thompson, Michael D.
TITLE OF INVENTION: Nucleic Acid Molecules And Oth
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(5193) B
CURRENT APPLICATION NUMBER: US/09/874,708A
CURRENT FILING DATE: 2001-12-20
PRIOR APPLICATION NUMBER: US 60/211,750
PRIOR FILING DATE: 2000-06-15
NUMBER OF SEQ ID NOS: 91742
SEQ ID NO 16025
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                                                                                                                                                          ; NAME/KEY: unsure
; LOCATION: (1)..(447)
; COTHER INFORMATION: unsure at all n locations
; OTHER INFORMATION: Clone ID: uC-gmrominsoy099e02b1
US-09-874-708A-16025
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US-09-874-708A-16025
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                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                   ENGTH: 447
                                                                                                    Local Similarity 85.2%;
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Similarity 85.2%;
            CAGTAGAACGAGCAAAACAAGAAGGAAATCTTCCAGTTTATGGCTACCA 233
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                                                                                  Score 153.8; DB 37; Length 447; Pred. No. 5.6e-37; 0; Mismatches 32; Indels 2;
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Pred. No. 4.5e-37;
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Search completed: May 4, Job time: 1836.74 secs
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                                                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: DNA
; ORGANISM: Glycine max
; OTHER INFORMATION: Clone ID: uC-gmrominsoy099e02b1
US-60-211-750-15882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-60-211-750-15882; Sequence 15882, Application US/60211750; GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/60/211,750
CURRENT FILING DATE: 2000-06-15
NUMBER OF SEQ ID NOS: 90609
SEQ ID NO 15882
                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 447
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                                                               CAGTAGAACGAGCAAAACAAGAAGGAAATCTTCCAGTTTATGGCTACCA 249
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2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

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GenCore version 5.1.6 (c) 1993 - 2005 Compugen Ltd.
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US-09-280-312D-618
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US-09-667-857-378
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3134, App
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19, Appl
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314, App
32950, A
              389, App
388, App
206, App
208, App
239, App
31, Appl
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 Query Match 17.3%; Best Local Similarity 50.0%; Matches 137; Conservative

Score 54.8; DB 4; Pred. No. 5.6e-09; 0; Mismatches 137;

Length :

0

RESULT 1 US-09-620-312D- Sequence 617, Patent No. 65 GENERAL INFORM APPLICANT: APPLICAN		C 28 33. 33. C 440 33. C 441 3 3. C 441 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4
2D-617 Application 1 656966 E56966 FORMATION: Tang, Y. Tom Liu, Chenghua Liu, Junding Liu, Yang, Yonghon Liu, Yang, Junrui Liu, John Tillingh Liu,		34 10.8 3693 3 .8 10.7 3639 2 .4 10.6 263693 4 .4 10.6 263693 4 .4 10.6 263694 4 .2 10.5 263594 4 .3 10.4 42574 4 31 10.4 1149 3 .2 10.1 1698 4 32 10.1 1998 4 32 10.1 2999 4 32 10.1 2995 4 33 10.4 41529 3 32 10.1 4403765 3 32 10.1 4403765 3 32 10.1 4403765 3
JS/09620312D JS/09620312D JS/09620312D JS/09620 Nucleic Acids and Deptides JS/09/620,312D JS/09/52,317 JS/09/52,317 JS/09/52,317 JS/09/52,317 JS/09/488,725 JS/09/488,725	ALIGNMENTS	US-09-056-226-1 US-08-737-524B-26 US-09-444-905-1 US-09-348-352-1855 US-09-949-016-16915 US-09-949-016-17525 US-09-949-016-17525 US-09-949-016-16515 US-09-257-583-4 US-09-250-739-67 US-09-949-016-16515 US-09-050-739-69 US-09-050-739-69 US-09-103-840A-2 US-09-103-840A-2 US-09-103-840A-1 US-09-103-840A-1 US-09-103-840A-1 US-09-949-001-760
		Sequence 1, Appli Sequence 26, Appl Sequence 1855, Appli Sequence 1855, Appli Sequence 18915, Appli Sequence 1, Appli Sequence 17525, Appli Sequence 4, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 69, Appli Sequence 16, Appli Sequence 16, Appli Sequence 16, Appli Sequence 2, Appli Sequence 16, Appli Sequence 16, Appli Sequence 17, Appli Sequence 1, Appli Sequence 1, Appli

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APPLICANT: "ANY TING
APPLICANT: Ma, Yunqing
APPLICANT: Ma, Yunqing
APPLICANT: Wang, Dunrui
APPLICANT: Wang, Dunrui
APPLICANT: Wang, Zhiwei
APPLICANT: John Tilinghast
APPLICANT: John Tilinghast
APPLICANT: Drmanac, Radoje T.
ITILE OF INVENTION: No. 6569662el Nucleic Acids and
ITILE OF INVENTION: No. 6569662el Nucleic Acids and
ITILE OF INVENTION: Polypeptides
FILE REFERENCE: 784CIP2B
CURRENT APPLICATION NUMBER: US/09/620,312D
CURRENT FILING DATE: 2000-07-19
FRIOR APPLICATION NUMBER: 09/552,317
PRIOR APPLICATION NUMBER: 09/488,725
PRIOR APPLICATION NUMBER: 09/488,725
PRIOR APPLICATION NUMBER: 09/488,725
PRIOR APPLICATION NUMBER: 09/488,725
PRIOR FILING DATE: 2000-01-21
NUMBER OF SEQ ID NOS: 1105
SOFTWARE: pt FL-genes Version 1.0
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; Sequence 618, Application US/09620312D
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ADDITCANT: Liu,
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TYPE: DNA
ORGANISM: Homo sapiens
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Ren, Feiyan
Chen, Rui-hong
Zhao, Qing A.
Wehrman, Tom
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Zhou, Ping
                                                                                                                                                                                                                                                                                                                                                                          Conservative
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idi, Vinod
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                                                                                                                                                                                                                                                                                                                                                                                                         17.3%;
                                                                                                                                                                                                                                                                                                                                                         Score 54.8; DB 4; Length 96
Pred. No. 5.8e-09;
0; Mismatches 137; Indels
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SOFTWARE: PERI
SEQ ID NO 3583
LENGTH: 271
TYPE: DNA
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; ORGANISM: Candida albicans
US-09-248-796A-3134
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                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                             APPLICANT: Lalgudi, Raghunath V.
APPLICANT: Ito, Laura Y.
APPLICANT: Ito, Laura Y.
APPLICANT: Ito, Laura Y.
APPLICANT: Sherman, Bradley K.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN EAR FILE REFERENCE: PL-0017 US
CURRENT APPLICATION NUMBER: US/09/313,294A
CURRENT FILING DATE: 1999-05-14
RUMBER OF SEQ ID NOS: 7600
SOFTWARE: PERL PROGRAM
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LENGTH: 795
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GENERAL INFORMATION:
APPLICANT: Keith We
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3134, Application US/09248796A Patent No. 6747137
FRATURE:
NAME/KEY: misc feature
OTHER INFORMATION: Incyte ID No.
NAME/KEY: unsure
LOCATION: 244, 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/248,796A CURRENT FILING DATE: 1999-02-12 PRIOR APPLICATION NUMBER: US 60/074,725 PRIOR FILING DATE: 1998-02-13 PRIOR APPLICATION NUMBER: US 60/096,409 PRIOR FILING DATE: 1998-08-13 PRIOR FILING DATE: 1998-08-13 PRIOR FILING DATE: 1998-08-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Keith Weinstock et al TITLE OF INVENTION: NUCLEIC ACID AM AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: 107196.132
                                                                                                           ORGANISM: Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  184
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            709 GCTATTATGAGGAGTGAAGACCCCAGATCTGTGATCAATCTATTAAGAAATGTTTGCTCA 768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 AAGGTGCGCACACTGAGAAGAAGTACCCTTCCCTTGACATAGAGGTTGATGGTGGTCGTA 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAGGTCGAAACATTAAGAGAAAAGTACCCCAACTTGGATGTTCAAGTTGATGGAGGATTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TCGGTGTTTAAACCAGAAGACCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGGAAGGATACAATTCAACCTGCTGCAGATGCTGGTGCAAATGTCATTGTTGCTGGGACC 708
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CATGAAATCTCCCTTTTACTGGAAAACAGGAATA 862
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ilarity 60.1%;
Conservative
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Pred. No. 6.1e-08;
0; Mismatches 57;
                                           6476212 700611933H1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            731
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US-09-691-270A-19/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; FEATURE:
; NAME/KEY: unsure
; LOCATION: (367)
; OTHER INFORMATION: n = a,
US-09-691-270A-19
                                                                                                                                                                                                                                                                                                               RESULT 6
                                                                                                                                                                        Sequence 2422, Application US/09107532A
Patent No. 6583275
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm
TITLE OF INVENTION: NUCLEIC ACID
ENTEROCCOCCUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19, Application US/09691270A Patent No. 6627795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 6627795
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity Matches 71; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: PCT/US99/08789
PRIOR FILING DATE: 1999-04-22
NUMBER OF SEQ ID NOS: 28
SOFTWARE: MICROSOFT OFFICE 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/691,270A
CURRENT FILING DATE: 2000-10-18
PRIOR APPLICATION NUMBER: 60/083,042
PRIOR FILING DATE: 1998-04-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Williams, Mark E.
APPLICANT: Sean J. COUGhlan
TITLE OF INVENTION: CAROTENOID BIOSYNTHESIS ENZYMES
FILE REFERENCE: BB1115 US NA2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 1246
TYPE: DNA
ORGANISM: Zea mays
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                   1104 TTTTTTCATTTGTCACGTTTTAGTTTAAAATTGAACTAGCAGACGACAAATATTCGAGAA 1045
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                          1044 CGAGTGTATCCTTTC 1030
                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME THERAPEUTICS CORPORATION
                                                                                                                                                        NUMBER OF SEQUENCES: 7310
                                                                                                                                                                                                                                                                                                                                                                                                              300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         284 GACTGATATTCGAGAATGGAGGGAGTACTT 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      224 AAGTACTCCCTCCGTTTTTTTTATTCGTCGCGTTTTAGTTCAAACATGAACTAGCGGAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
57; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                              TGGAGGGAGTACTTC 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TTTTTTTATTCGTCGCGTTTTAGTTCAAACATGAACTAGCGGACGACTGATATTCGAGAA 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGCAAATATTCGAGAACAGATGTAGTACTT 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAATAATCCCNCCATTCTTTTTATNTGTCGCGGTTTAGTTTGAA-ATGAACTAGCGGGC
                                                        STATE: Massachusetts
                                      COUNTRY: USA
                                                                             CITY: Waltham
                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                               100 Beaver Street
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Pred. No. 7.9e-06;
0; Mismatches 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 50.4; DB 4; Pred. No. 1.1e-07;
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                                                                                                                                                                        and David Bush
AND AMINO ACID SEQUENCES RELATING TO
FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 1246;
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; LOCATION: (B) LOCATION 1...663;
; SEQUENCE DESCRIPTION: SEQ ID NO: 2422:
US-09-107-532A-2422
                                                                                                                                                                                                                                                                                                                                                                            RESULT 7
; TYPE: DNA
; ORGANISM: Homo
US-09-404-879A-378
                                                                                                                                                                                                                                                                                                                                                     US-09-404-879A-378/C
                                                        CURRENT APPLICATION NUMBER: US/09/404,879A
CURRENT FILING DATE: 1999-09-24
NUMBER OF SEQ ID NOS: 393
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 378
LENGTH: 395
                                                                                                                                                                                                                                                                                                             Sequence 378, Application US/09404879A Patent No. 6468546
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                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                        APPLICANT: Algate, Paul A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.462C2
                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 2422: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 12.0%;
Local Similarity 55.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
TELEPHONE: (781)893-5007
TELEFAX: (781)893-8277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/051571
FILING DATE: July 2, 1997
ATTORNEY/ARGENT INFORMATION:
NAME: Ariniello, Pamela Denek
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA: US/09/107,532A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              552
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                                                                                                                                                                                                                                                                                                                                                                                                                                   612 AGCCGAAAATCCAG 625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            75 CATAGACGIGGCCGCATCIGCGGGCCAAITGCAICGICGCIGGAAGCICTAIAIIIIGG 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15 ACTGAGAAAGAAGTACCCTTCCCTTGACATAGAGGTTGATGGTGGTCTAGGTCCTTCAAC 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        74;
                                                                                                                                                                                                                                                    Mitcham, Jennifer L. King, Gordon E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AGCAAAATCTGTAAAGATGCAGGGGGGGGGATGTCTTCGTTGCAGGATCATATATCTATGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACGTAGAGAGAACAATTGGACTTATGCAATCGAAGTAGATGGCGGAATCGTTCCAGAAAC 551
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: PC
OPERATING SYSTEM: <Unknown>
SOFTWARE: ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CGCTGCGGACCCAG 148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Enterococcus faecium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 60/085,598 FILING DATE: 14 May 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 30-Jun-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: CD/ROM ISO9660
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>.</u>.
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Pred. No. 0.0047;
0; Mismatches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 663;
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611

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APPLICANT: Algate, Paul A.

APPLICANT: Fling, Steven P.

APPLICANT: Fetter, Marc W.

APPLICANT: Fanger, Gary Richard

APPLICANT: Feed, Steven G.

APPLICANT: Vedvick, Thomas S.

APPLICANT: Vedvick, Thomas S.

APPLICANT: Vedvick, Thomas S.

ITITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

ITITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.462C5

CURRENT APPLICATION NUMBER: US/09/667,857

CURRENT APPLICATION NUMBER: US/09-20

NUMBER OF SEQ ID NOS: 455

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 378

LENGTH: 395
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; ORGANISM: Homo sapiens
US-09-667-857-378
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                                                                                                                                                                                                                                                                                               Sequence 314, Application US/09854133 Patent No. 6759508
SEQ ID NO 314
                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
      APPLICANT: Henderson, Robert A.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: THE THERAPY AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.475C10
CURRENT APPLICATION NUMBER: 10/09/854,133
CURRENT FILING DATE: 2001-05-11
NUMBER OF SEQ ID NOS: 735
SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
                                                                                                                                                                 APPLICANT: Lodes, Michael J.
APPLICANT: Mohamath, Raodoh
APPLICANT: Henderson, Robert
APPLICANT: Benson, Darin R.
APPLICANT: Secrist, Heather
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Best Local Similarity 63.1%;
Matches 53; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 117 TGGAAGCTCTATATTTGGCGCTGC 140
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
SEQIDNO 22949
LENGTH: 601
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US-09-949-016-22950/c
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; ORGANISM: Human
US-09-949-016-22949
; Sequence 22950, Application US/09949016
; Patent NO. 6812339
; GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: MITH HUMAN DISBASE, METHODS OF DETECTION AND USES THEREOF
TILE REFERENCE: CL001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT APPLICATION NUMBER: 00/241,755
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR APPLICATION NUMBER: 60/237,768
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; NAME/KEY: misc_feature
; LOCATION: (1)...(591)
; OTHER INFORMATION: n = A,T,C or
US-09-854-133-314
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GENERAL INFORMATION:
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Best Local Similarity
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                        117
                                                                                                                                                                                                                                                                                                                                                                                                        164 TGGTGCAGGTGCTAACAGTAGGAGTGGCTGGTTCTGGTGCTGGTATTGGAACAGTCTT 105
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ilarity 63.1%;
Conservative
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Pred. No. 0.086;
0; Mismatches 31; Indels
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Pred. No. 0.086;
0; Mismatches 31; Indels
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31;
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; LENGTH: 601
; TYPE: DNA
; ORGANISM: Human
US-09-949-016-22951
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; ORGANISM: Human
US-09-949-016-22950
                                        Sequence 195227, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION: APPLICANT: VENTER, J.
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Patent No. 6812339
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
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PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FASTSEQ for Windows Version 4.0
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)EQ ID NO 22950
LENGTH: 601
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PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
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63.1%;
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Pred. No. 0.086;
0; Mismatches 31;
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                                         APPLICANT: VENTER, J.
                                                                               Sequence 195229, Application US/09949016 Patent No. 6812339
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Best Local :
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
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PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
     TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
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ORGANISM: Human
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J. Craig et al.
POLYMORPHISMS IN KNOWN GENES
DISEASE, METHODS
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Pred. No. 0.086;
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FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR PPLICATION WIMBER: 60/241,755

PRIOR PPLICATION WIMBER: 60/237,768

PRIOR APPLICATION WIMBER: 60/237,768

PRIOR APPLICATION WIMBER: 60/231,498

PRIOR APPLICATION WIMBER: 60/231,498

PRIOR PILING DATE: 2000-10-03

PRIOR APPLICATION WIMBER: 60/231,498

PRIOR PILING DATE: 2000-09-08

INDIA DATE: 2000-10-03

PRIOR PILING DATE: 2
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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                 4 US-09-313-294A-2427

4 US-09-620-312D-617

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4 US-09-620-312D-618

4 US-09-620-312D-618

4 US-09-620-312D-618

4 US-09-183-110-947

4 US-09-107-433-453

4 US-09-107-433-453

4 US-09-1107-532A-242

2 US-09-1107-532A-242

4 US-09-1107-532A-242

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10 4 US-09-138-185A-1

10 4 US-09-231-0178-152

11 US-09-231-0178-152

12 US-09-218-152-10178-152

13 US-09-218-152-10178-152

14 US-09-710-279-307

15 US-09-710-279-307

17 US-09-710-279-3257

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              Sequence 2427, Ap Sequence 618, Appl Sequence 1149, Ap Sequence 1149, Ap Sequence 143, App Sequence 1299, Ap Sequence 1299, Ap Sequence 149, Appl Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1080, Ap Sequence 1080, Ap Sequence 2557, Ap Sequence 2557, Ap Sequence 3071, Ap Sequence 3071, Ap Sequence 3071, Ap Sequence 3073, Ap Sequence 3134, Ap Sequence 3
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CTGGTG	CTTCAC AGACAI	nilarit Conse TCCACA TACATA	294A-2427 294A-2427 information: 6.6476212 information: MT: Lalgudi, Ra MT: Ito, Laura NT: Sherman, Br FINVENTION: PO PERENCE: PL-001 APPLICATION NUM FILLING DATE: 1 OF SEQ ID NOS: E: PERL PROGRAMME E: 267 DNA C 2427 DNA SM: Zea mays E: 267 DNA E: 267 DNA C 2427 DNA E: 165 feature Information: Inc	13.6 13.2 13.2 13.2 12.6 11.9	
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71 .	TCCTCTTGATTATGTTGAACCCTTGGCAAAAGCTGGTGCTTCTGGTTTTACATTTCACG	40.0%; Score 155.2; DB 4; Length 267; imilarity 75.4%; Pred. No. 1.5e-42; Conservative 0; Mismatches 63; Indels 0; Gaps GCTCCACATGGACTACTGGATGGGCATTTTGTCCCCACATTTAACTATTGGGGCTCCAG GCTACATATGGACTACTGGGGCTCCGGGCTACATATTGGACTATTGGACTATTGGGCTCCCGGTATATTGGACATATTGGACTATTGGACTATTGGACTATAGATACTAAAGTTTGAAAAGTACAAAAGACAAAAGAAATTTGGATTACAATATTTGGAAAGTTTGAAAAGTAAACATACCAAAGCATATTTGGACTGCCATCTTATGGTTACAATATTGGACTGCCATCTTATGGTCACAAAGCATAATTTGGACTGCCATCTTATGGTCACAAAGCATATTTGGACTGCCATCTTATGGTCACAAAGCATATTTGGACTGCCATCTTATGGTCACAAAGCATATTTGGACTGCCATCTTATGGTCACAAAGCATATTTGGACTGCCATCTTATGGTCACAAAGCATATTTGGACTGCCATCTTATGGTCACAAAGCATATTTGGACTGCCATCTTATGGTCACAAAGCATATTTGGACTGCCATCTTATGGTCACAAA	tion US/09313294A Raghunath V. a Y. Bradley K. POLYNUCLEOTIDES AND POLYPEPTIDES 017 US UMBER: US/09/313,294A 1999-05-14 7600 am	US-09-643-990A-1 US-09-513-999C-1816 US-08-916-421B-1 US-09-692-570-1 US-09-596-002-31 US-09-790-988-31 US-09-252-991A-825 US-09-489-039A-2525 ALIGNMENTS	US-09-103-840A-1 US-08-714-918-79 US-09-265-315-79 US-09-265-315-79 US-09-266-417-79 US-09-528-709-79 US-09-527-745-79 US-09-489-039A-2344 US-09-489-039A-2344
	TTACATTTCACGT 29	### A0.0%; Score 155.2; DB 4; Length 267; imilarity 75.4%; Pred. No. 1.5e-42; Conservative 0; Mismatches 63; Indels 0; Gaps 0; GCTCCACATGGACATCATGGATGGGCATTTTGTCCCCAATTTAACTATTGGGCGCTCCAGT 175 GCTACATATGGACATCATGGATGGGCACTTCTGTCCTAACCTGATGGGGCTCCCGGT 60 GCTACATATGGACATCATGGATGGATGGATGGACATATTTGGACTATTGGGGCTTCCGGT 60 TATTGAAAGTTTGAGAAAGCACAAAGCATACTTTGGATTGTCACCTTATGGTTACAAA 235	DES DERIVED FROM CORN EAR	Sequence 1, Appli Sequence 1816, Ap Sequence 1, Appli Sequence 1, Appli Sequence 31, Appli Sequence 31, Appli Sequence 1, Appli Sequence 255, App Sequence 2525, App	Sequence 1, Appli Sequence 79, Appl Sequence 79, Appl Sequence 79, Appl Sequence 79, Appl Sequence 79, Appl Sequence 79, Appl Sequence 2344, Ap Sequence 1855, Ap Sequence 1857, Appli

RESULT 2 US-09-620-312D-617

Sequence 617, Application US/09620312D Patent No. 6569662

INFORMATION:

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APPLICANT: Drmanac, Radoje T.

ITILE OF INVENTION: No. 656962el Nucleic Acids and ITILE OF INVENTION: Polypeptides

FILE REFERENCE: 784CIP2B

CURRENT APPLICATION NUMBER: US/09/620,312D

CURRENT FILING DATE: 2000-07-19

PRIOR APPLICATION NUMBER: 09/552,317

PRIOR FILING DATE: 2000-04-25

PRIOR APPLICATION NUMBER: 09/488,725

PRIOR APPLICATION NUMBER: 09/488,725

PRIOR FILING DATE: 2000-01-21

NUMBER OF SEQ ID NOS: 1105

SOFTWARE: pt FL genes Version 1.0

SEQ ID NO 617
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APPLICANT: Liu, Chenghua
APPLICANT: Asundi, Vinod
APPLICANT: Zhang, Jie
APPLICANT: Ren, Feiyan
APPLICANT: Chen, Rui-hong
APPLICANT: Zhao, Qing A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (61)..(747)
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433 GT 434
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                            GT 388
                                                    ATTAAAGACATTCGGGAGAATGGGATGAAGGTTGGCCTTGCCATCAAACCAGGAACCTCA 432
                                                                                 GCTGGTGCTTCTGGTTTTACATTTCACGTAGAGACATCAAAAGATAACTGGAAAGAACTT
                                                                                                                   GCAGGAGCCAATCAGTACACCTTTCATCTCGAG---GCTACTGAGAACCCCAGGGGCTTTG
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Wang, Dunrui
Wang, Zhiwei
John Tillinghast
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Xue, Aidong J.
Yang, Yonghong
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Zhou, Ping
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56.9%;
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Pred. No. 1.2e-19;
0; Mismatches 147; Indels
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US-09-276-531-8
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Best Local Similarity
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Patent No.
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APPLICANT: Bandman
APPLICANT: Lal, pr
APPLICANT: Hilman
APPLICANT: Yue, He
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (650) 845-410 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Lynn E. Murry, Ph.D.
REGISTRATION NUMBER: 42,918
REFERENCE/DOCKET NUMBER: PA-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 855-0555
                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linea
IMMEDIATE SOURCE:
LIBRARY: LIVRTU
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 6:
FILING DATE: March 27:
CLASSIFICATION:
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LENGTH: 3556 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/276,531
FILING DATE: Herewith
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 1
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APPLICANT: Baughn, Mariah R.
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF GENES
TITLE OF INVENTION: RECEPTORS AND PROTEINS ASSOCIATED WITH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                CLONE:
                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: single
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        252
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                                                                                                                                                33 AAAATAGCTCCTTCGATGCTCTCTCCGACTTCGCCAATTTGGCTTCCGAGGCCTCAGCGC 92
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5. 6183968
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PALO ALTO
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TTCTTTGACATGCACATGATGGTGTCCAAGCCAGAACAGTGGGTAAAGCCCAATGGCTGTA
                           TATTTGGATTGTCACCTTATGGTTACAAATCCTCTTGATTATGTTGAACCCTTGGCAAAA 266
                                                           AACATCACCTTTGGTCACCCTGTGGTAGAAAGCCTTCGAAAGCAGCTAGGCCAGGACCCT
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(650) 8:
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Hillman, Jennifer
Yue, Henry
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Pred. No. 2.7e-19;
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RESULT 4
US-09-620-312D-618
; Sequence 618, Application US/09620312D
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; LOCATION: (61)..(801)
US-09-620-312D-618
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PRIOR FILING DATE: 2000-04-25
PRIOR APPLICATION NUMBER: 09/488,725
PRIOR FILING DATE: 2000-01-21
NUMBER OF SEQ ID NOS: 1105
SOFTWARE: pt_FL_genes Version 1.0
SEQ ID NO 618
                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                            Query Match
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APPLICANT:
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TITLE OF INVENTION: No. 6569662el Nucleic Acids and
TITLE OF INVENTION: Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: 784CIP2B
CURRENT APPLICATION NUMBER: US/09/620,312D
CURRENT FILING DATE: 2000-07-19
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                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                   ENGTH: 961
                                                                                                                                                                                                                                              Local Similarity
                 207
                                                                                                         136
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Ren, Feiyan
Chen, Rui-hong
Zhao, Qing A.
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TATTTGGATTGTCACCTTATGGTTACAAATCCTCTTGATATGTTGAACCCTTTGGCAAAA 266
                                                                          AATTTAACTATTGGCGCTCCAGTTATTGAAAGTTTGAGAAAGCA-----CACAAAGGCA 206
                                                                                                         ATGCTAGACTCTGGGGCCGATTATCTGCACCTGGACGTAATGGACGGGCATTTTGTTCCC
                                                                                                                                    AAGATTGGCCCGTCCATCCTCAACAGCGACCTGGCCAATTTAGGGGCCGAGTGCCTCCGG 135
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Wang, Dunrui
Wang, Zhiwei
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                                             AACATCACCTTTGGTCACCCTGTGGTAGAAAGCCTTCGAAAGCAGCTAGGCCAGGACCCT
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Asundi, Vinod
Zhang, Jie
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John Tillinghast
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Yang, Yonghong
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                                                                                                                                                                                                                                             21.9%;
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                                                                                                                                                                                                                                             Score 85; DB 4;
Pred. No. 2.4e-18;
                                                                                                                                                                                                                                Mismatches 105; Indels
                                                                                                                                                                                                                                                           Length 961;
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                                                                                                                                                                                                                             Gaps
                                                                                                        195
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; SEQUENCE DESCRIPTION: SEQ ID NO: 1149: US-08-956-171E-1149
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US-08-956-171E-1149/c
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (240) 314-1224
TELEFAX: (301) 309-8439
INFORMATION FOR SEQ ID NO: 1149:
                                                                                                                                                                                                                                                                                                            Match 21.5%;
Local Similarity 53.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: January 3, 1997
ATTORNEY/AGENT INFORMATION:
NAME: MARK J. Hyman
REGISTRATION NUMBER: 46,789
REFERENCE/DOCKET NUMBER: PB248P1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
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Michael R. Fannon
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
NUMBER OF SEQUENCES: 5256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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                                                                                           745 CAAAACTATATCCATCATTATCTGTTGATTTTTTGGATTTACAACATGAATTAAAAC 686
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                                                                                                                                 31 CGAAAATAGCTCCTTCGATGCTCTCTCCGACTTCGCCAATTTGGCTTCCGAGGCTCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 916 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 60/009,861 FILING DATE: January 5, 1996 APPLICATION NUMBER: 08/781,986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/956,171E FILING DATE: 20-Oct-1997 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Diskette, 3.50 inch, 1
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOPTWARE: ASCII Text
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 9410 Key West Avenue
CITY: Rockville
STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GCTGGTGCTTCTGGTTTTACATTTCACGTAGAGAC 301
                                        CCAATTTAACTATTGGCGCTCCAGTTATTGAAAGTTTGAGAAAGCACACAAAGGCATATT 210
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CTAATATATCTATTGGTTTACCAATATTAGATGCAGTAAGAAAAGGCACAACATTACCTA
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                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patrick S. Dillon
Craig A. Rosen
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                                                                                                                                                                                                                                                                                       Score 83.6; DB 4;
Pred. No. 7.1e-18;
2; Mismatches 161;
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US-08-781-986A-1149/c
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                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Benson, Bob
REGISTRATION NUMBER: 30,446
REFERENCE/DOCKET NUMBER: PB:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                      LENGTH: 916 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: MS
SOFTWARE: ASCII Text
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
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TGGATTGTCACCTTATGGTTACAAATCCTCTTGATTATGTTGAACCCTTGGCAAAAGCTG
                                                                   CCAATTTAACTATTGGCGCTCCAGTTATTGAAAGTTTGAGAAAGCACACAAAGGCATATT
                                                                                                                                                                CAAAACTATATCCATCATTATTATCTGTTGATTTTTTGGATTTACAACATGAATTAAAAC
                                             CTAATATATCTATTGGTTTACCAATATTAGATGCAGTAAGAAAAGGCACAACATTACCTA
                                                                                                     gcargcrccacrrcgccccarragcrccacarcarcarcarcarcarcarcacacarrrrrgrcc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GTGCCGATATGATTTCAATTCATGTCGA---ATCAACGCCTCATATTCATCGTGCTATTC
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5255
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486/33
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Pred. No. 7.1e-18;
2; Mismatches 161;
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RESULT 8
US-09-107-433-453
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; LENGTH: 903;
; TYPE: DNA; ORGANISM: Streptococcus pneumoniae
US-09-583-110-947
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Best Local Similarity 53.7
191; Conservative
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Patent No. 6699703
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: PATHO0-07A

CURRENT APPLICATION NUMBER: US/09/583,110

CURRENT FILING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: US 09/107,433

PRIOR FILING DATE: 1998-06-30

PRIOR APPLICATION NUMBER: US 60/085,131

PRIOR FILING DATE: 1998-05-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Lynn Doucette-Stamm et al.
TITLE OF INVENTION: Nucleic Acid and Amino Acid Sequences Relating
TITLE OF INVENTION: Pneumoniae for Diagnostics and Therapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/051,553
PRIOR FILING DATE: 1997-07-02
NUMBER OF SEQ ID NOS: 5322
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                                                                                              AGAATCAAGTCACATGGCATGATTCCTGGTGTAGCATTAAAGCCCTGGGACCCCCGT 388
                                                                                                                                                                      GCTTCTGGTTTTACATTTCACGTAGAGACATCAAAAGATAACTGGAAAGAACTTATCCAA 332
                                                                                                                                                                                                          GATTGCCACTTGATGGTGTCAAACCCTGAGCATCATCTGGAAGATTTTGCGCGTGCAGGT
                                                                                                                                                                                                                                      GATTGTCACCTTATGGTTACAAATCCTCTTGATTATGTTGAACCCCTTGGCAAAAGCTGGT 272
                                                                                                                                                                                                                                                                            CAAATCAGTTTTGGTGCAGGTGTGGTCGAGAGCCTTCGTCCTCATAGTAAGATGGTTTTC 438
                                                                                                                                                                                                                                                                                                                AAITTAACTATIGGCGCTCCAGTTATTGAAAGTTTGAGAAAGCACACAAAAGGCATATTTG 212
                                                                                                                                                                                                                                                                                                                                                                                    AAAATTCGTTCACTCGGAGTTAAGCCTTCAGTCGTTATCAATCCTGGCACACCAGT
                                                                                                                                    GCAGACATCATCAGTATCCATGTAGA---AGCAACACCTCATATTCATGGCGCCCTCCAA
                                                                                                                                                                                                                                                                                                                                                    CTAGAAGCAACTGGGGCAGAATATGCCCATATCGATATCATGGACAGTCATTTTGTACCG
                                                                                                                                                                                                                                                                                                                                                                                                                        AAGATTGCTCCGTCAATTCTGGCAGCAGATTATGCCAACTTTGAACGTGAAATCAAACGT 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAAATAGCTCCTTCGATGCTCTCTTCCGACTTCGCCAATTTTGGCTTCCGAGGCTCAGCGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20.8%;
53.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches 162;
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Pred. No. 6.3e-17;
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Sequence 453, Application US/09107433

Patent No. 6800744

GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID
SEQUENCES RELATING TO STREPTOCCCCUS
THERAPEUTICS

PNEUMONIAE

FOR

DIAGNOS

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; NAME/KEY: misc_feature; LOCATION: (B) LOCATION 1...993; SEQUENCE DESCRIPTION: SEQ ID NO: 453: US-09-107-433-453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 191; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/085131

FILING DATE: May 12, 1998
APPLICATION NUMBER: 60/05153

FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Ariniello, pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: «Unknown»
CURRENT APPLICATION NUMBER: US/09/107,433
APPLICATION NUMBER: US/09/107,433
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MEDIUM TYPE: CD/ROM ISO9660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL: N
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646
                                                                                                                                                                                       529
                                                                                                                                                                                                                                                                                   469
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                                                                                                                                                                                                                                                                                                                             153 AATTTAACTATTGGCGCTCCAGTTATTGAAAGTTTGAGAAAGCACACAAAGGCATATTTG 212
                                                                                                                                                                                                                                                                                                                                                                               409
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                                                                                                                                                                                                                                                                                 CAAATCAGTTTTGGTGCAGGTGTGGTCGAGAGCCTTCGTCCTCATAGTAAGATGGTTTTC
                                                                                                                                   GCTTCTGGTTTTACATTTCACGTAGAGACATCAAAAGATAACTGGAAAGAACTTATCCAA 332
                                                                                                                                                                                       GATTGCCACTTGATGGTGTCAAACCCTGAGCATCATCTGGAAGATTTTGCGCGTGCAGGT
                                                                                                                                                                                                                                                                                                                                                                               CTAGAAGCAACTGGGGCAGAATATGCCCATATCGATATCATGGACAGTCATTTTGTACCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 993 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: <Unknown>
OPERATING SYSTEM: <Unknown>
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STATE: Massachusetts
AAAATTCGTTCACTCGGAGTTAAGCCTTCAGTCGTTATCAATCCTGGCACACCAGT 701
                                          AGAATCAAGTCACATGGCATGATTCCTGGTGTAGCATTAAAGCCTGGGACCCCCGT 388
                                                                                         GCAGACATCATCAGTATCCATGTAGA----AGCAACACCTCATATTCATGGCGCCCTCCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (781)893-5007
TELEFAX: (781)893-8277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
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Pred. No. 6.7e-17;
0; Mismatches 162;
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NAME/KEY: misc_feature;
COATION: (B) LOCATION 1...663;
SEQUENCE DESCRIPTION: SEQ ID NO: 2422;
US-09-107-532A-2422
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US-09-107-532A-2422
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Patent No. 6583275
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm
TITLE OF INVENTION: NUCLEIC ACID
ENTEROCCOCCUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Ariniello, Pamela Deneke REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
TELEPHONE: (781)893-5007
TELEFAX: (781)893-8277
INFORMATION FOR SEQ ID NO: 2422:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 195; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: PC
COMPUTER: PC
OPERATING SYSTEM: «Unknown»
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/09/107,532A
FILING DATE: 30-Jun-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/085,598
FILING DATE: 14 May 1998
APPLICATION NUMBER: 60/051571
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: circular MOLECULE TYPE: DNA (gend HYPOTHETICAL: NO ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
COMPUTER: PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
183
                                          200
                                                                                      123
                                                                                                                                                                                                                         80
                                                                                                                                                                             20 GGGAATGACACCGAAAATAGCTCCTTCGATGCTCTTCCGACTTCGCCAATTTGGCTTC
                                                                                                                                                                                                                                                                   3 GGAAACAAGAATGAAGATAGCACCATCCATTTTGAGTGCAGATTTTGCCAATTTTACAAAG 62
                                                                                                             GCATTTTGTCCCCAATTTAACTATTGGCGCTCCAGTTATTGAAAGTTTGAGAAAGCACAC 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Waltham
STATE: Massachusetts
TAAATTACCGTTAGACGTCCATCTGATGATCGTTGATCCAGAAAACTACATTCCAGCATT
                                          AAAGGCATATTTGGATTGTCACCTTATGGTTACAAATCCTCTTGATTATGTTGAACCCCTT 259
                                                                                      CCAATTTGTCCCAAATATCACATTAGGGCCAAATATTGTTCAGGCAATCCGTCCTGTTAC
                                                                                                                                                                                                                  TYPE: nucleic acid STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Enterococcus faecium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: GENOME THERAPEUTICS CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENGTH: 663 base pairs,
                                                                                                                                                                                                                                                                                                                                                          20.5%;
llarity 52.8%;
Conservative
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Pred. No. 1.6e-16;
0; Mismatches 171;
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AND AMINO ACID SEQUENCES RELATING TO
FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                      Length 663;
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APPLICANT: Lynn Doucette-Stamm et al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACITILE OF INVENTION: ENTEROCOCCUS FAECALIS FOI FILE REFERENCE: 032796-032
CURRENT APPLICATION NUMBER: US/09/134,000C
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/055,778
PRIOR FILING DATE: 1997-08-15
NUMBER OF SEQ ID NOS: 6812
SOFTWARE: Patentin version 3.1
LEUGTH: 690
                                                  RESULT 11
US-08-961-527-149
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Sequence 149, Application US/08961527 Patent No. 6420135 GENERAL INFORMATION:
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US-09-134-000C-1299
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Enterococcus faecalis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9 AAAGAAAAGATGGGAATGACGAAAATAGCTCCTTCGATGCTCTCTTCCGACTTCGCC 68
                                                                                                                                                                                                                                                                    GŢŢĠĄĀĊĊŢŢĠĊĀĀĀĀĠĊŢĠŢĠŢŢĊŢŢĊĠŢŢŢŢĀĊĀŢŢĊĀĊĠŢĀĠĀĠĀĊĀŢĊĀĀĀĀ 308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                               TTAPAGCCTGGGACCCC 385
                                                                                                                                                                         CATA---TTCATCGTGCCTTACAAATGATTAAAAATGCGGGTGTGAAAGCTGGCGTAACC
                                                                                                                                                                                                   GATAACTGGAAAGAACTTATCCAAAGAATCAAGTCACATGGCATGATTCCTGGTGTAAGCA 368
                                                                                                                                                                                                                                        ATCAATGCTTTTGCGGAAGCCGGTGCGGATATTATTACCGTCCATCAAGAAGCAACACCC 315
                                                                                                                                                                                                                                                                                                       CGTCCAGTGÁCAAAACTÁCCATTGGÁTGTTCÁTTTGÁTGATTGTÁCAACCTGAAAATTÁT 255
                                                                                                                                                                                                                                                                                                                                    AGAAAGCACACAAAAGGCATATTTGGATTGTCACCTTATGGTTACAAATCCTCTTGATTAT 248
                                                                                                                                                                                                                                                                                                                                                                     GTCATGGACGGTCAATTTGTGCCCCAACATTACCTTAGGACCGAATATCGTTTCAGCGATT 195
                                                                                                                                                                                                                                                                                                                                                                                                   ATÇAŢĢĢĀTĢĢĢÇĀTŢŢŢĢŢĊÇÇĀATTŢĀĀÇTAŢTĢĢCGÇTCCAGŢTAŢŢĢĀĀAGTTŢG 188
                                                                                                                                                                                                                                                                                                                                                                                                                                      AATTTAGAAAGAGATATCCGTTTAGTAGAAGAATTAGGAGCAGATTATATTCATGTTGAT 135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AATTTGGCTTCCGAGGCTCAGCGCATGCTCCACTTCGGCGCGGATTGGCTCCACATGGAC 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AATATÄÄÄĠGGAĠCGĀĠCCAĀATĠĀĀĀĊTĀĠĊAĊĊĀŤĊĊĀŤTTĪAAGŤĠĊĊĠĀŦŤŤTĠĊĊ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TĠĊĀĀĀĀĠĊŤĠĠĀĠĊĀGĀŤĀTCĀŤŤĀĊTGŤĊĊĀĊĠŤĊĠĀĀĢĊĀĀĊTCCTCĀŤĀTTCĀTCG 302
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Pred. No. 5
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5.7e-16;
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FOR DIAGNOSTICS AND THER
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RESULT 12
US-09-198-452A-1/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: MSDOS
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Brookes, A. Anders
REGISTRATION NUMBER: 36,373
REFERENCE/DOCKET NUMBER: PB:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
COMPUTER: HP Vectra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Charles TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 9410 Key West Avenue CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Human Genome Sciences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                5026
                                                                                                                                                                                            4966
                                                                                                                                                                                                                                              4906 CAAATCAGTTTTGGTGCAGGTGTGGTCGAGAGCCTTCGTCCTCATAGTAAGATGGTTTTC
                                                                                                                                                                                                                                                                                                                      4846
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                                                                                                                                                                273
                                                                                                                                                                                                         213 GATTGTCACCTTATGGTTACGAATCCTCTTGATTATGTTGAACCCTTGGCAAAAGCTGGT 272
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                                                                                                                                                                                                                                                                                                                                                     93
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                          AAAATTCGTTCACTCGGAGTTAAGCCCTTCAGTCGTTATCAATCCTGGCACATCAGT 5138
                                                                                      AGAATCAAGTCACATGGCATGATTCCTGGTGTAGCATTAAAGCCCTGGGACCCCCGT 388
                                                                                                                                               GCTTCTGGTTTTACATTTCACGTAGAGACATCAAAAGATAACTGGAAAGAACTTATCCAA 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      I: 12566 base pairs
nucleic acid
EDNESS: double
                                                                                                                      GCAGACATCATCAGTATCCATGTAGA---AGCAACGCCTCATATTCATGGCGCCCCTCCAA
                                                                                                                                                                                    GATTGCCACTTGATGGTGTCAAACCCTGAGCATCATCTGGAAGATTTTGCGCGTGCAGGT
                                                                                                                                                                                                                                                                                                            CTAGAAGCAACTGGGGCAGAATATGCCCCATATCGATATCATGGACAGTCATTTTGTACCG 4905
                                                                                                                                                                                                                                                                              AATTTAACTATTGGCGCTCCAGTTATTGAAAGTTTGAGAAAGCACACAAAAGGCATATTTG
                                                                                                                                                                                                                                                                                                                                                                         AAGATTGCTCCGTCAATTCTGGCAGCAGATTATGCCAACTTTGAACGTGAAATCAAACGT
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                                                                                                                                                                                                                                                                                                                                 Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                           20.0%;
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Pred. No. 3.7e-15;
0; Mismatches 164;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PB340P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 12566;
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GENERAL INFORMATION: APPLICANT: Griffais

CANT: Griffais, OF INVENTION:

Chlamydia

pneumoniae genomic sequence and polypeptides, fragments

Sequence 1, Application US/09198452A Patent No. 6559294

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SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: thereof and TITLE OF INVENTION: and treatme FILE REFERENCE: 9710-003-999 CURRENT APPLICATION NUMBER: US/O CURRENT FILING DATE: 1998-11-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 1230025
TYPE: DNA
ORGANISM: Chlamydia pneumoniae
                                                                                                    LOCATION: (255001)..(270000)

OTHER INFORMATION: n=a or c or
                                                                                                                                               COCATION: (240001)..(255000)

DTHER INFORMATION: n=a or c or g or t
                            AME/KEY: misc_feature
OCATION: (285001)..(3
                                                  AME/KEY: misc feature
OCATION: (270001)..(285000)
WHER INFORMATION: n=a or c or
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OCATION: (1
                                                                                                                  AME/KEY: misc_feature
OCATION: (255001)..(270000)
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OCATION: (225001)..(240000)
THER INFORMATION: n=a or c or g
AME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                AME/KEY: misc_feature
OCATION: (180001)..(195000)
THER INFORMATION: n=a or c or g
                                                                                                                                                                                                                                        CHER INFORMATION: n=a or c
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CHER INFORMATION: n=a or c
AME/KEY: misc_feature
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THER INFORMATION: n=a or c or
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CHER INFORMATION: n=a or c or g
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CHER INFORMATION: n=a or c or g or t
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HER INFORMATION: n=a or c
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HER INFORMATION: n=a or c
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NAME/KEY: misc feature LOCATION: (615001)..(630000) OTHER INFORMATION: n=a or c OTHER INFORMATION: n=a or c or NAME/KEY: misc feature LOCATION: (63001)..(645000)
OTHER INFORMATION: n=a or c
NAME/KEY: misc_feature
LOCATION: (645001)..(660000) LOCATION: (600001)..(615000)
OTHER INFORMATION: n=a or c or NAME/KEY: misc_feature LOCATION: (570001)..(585000) OTHER INFORMATION: n=a or c or g or t LOCATION: (465001)..(480000) OTHER INFORMATION: n=a or c LOCATION: (435001)..(450000)
OTHER INFORMATION: n=a or c LOCATION: (390001)..(405000)
OTHER INFORMATION: n=a or c OTHER INFORMATION: n=a or c NAME/KEY: misc NAME/KEY: misc LOCATION: (585) LOCATION: (555001)..(570000)
DTHER INFORMATION: n=a or c or LOCATION: (360001)..(375000)
OTHER INFORMATION: n=a or c or COCATION: (330001)..(345000)
OTHER INFORMATION: n=a or c or g NAME/KEY: m: OTHER INFORMATION: n=a or c VAME/KEY: misc LOCATION: (5400 OCATION: (525001)..(540000)
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LOCATION: (73501)..(75000)
OTHER INFORMATION: n=a or c or g
NAME/KEY: misc feature
LOCATION: (75001)..(76500)
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NAME/KEY: misc feature
LOCATION: (765001)..(78000)
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NAME/KEY: misc feature
LOCATION: (780001)..(79500)
OTHER INFORMATION: n=a or c or g
NAME/KEY: misc feature
LOCATION: (79501)..(81000)
OTHER INFORMATION: n=a or c or g
NAME/KEY: misc feature
LOCATION: (810001)..(815000)
OTHER INFORMATION: n=a or c or g
NAME/KEY: misc feature
LOCATION: (82501)..(84000)
OTHER INFORMATION: n=a or c or g
NAME/KEY: misc feature
LOCATION: (82501)..(84000)
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LOCATION: (85501)..(870000)
OTHER INFORMATION: n=a or c or g or
NAME/KEY: misc_feature
LOCATION: (870001)..(885000)
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LOCATION: (900001)..(915000)
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LOCATION: (885001)..(900000)
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LOCATION: (705001)..(720000)
OTHER INFORMATION: n=a or c
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OTHER INFORMATION: n=a or c
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                              TTCTGGTTTTACATTTCACGTAGAGAÇATCAAAAGATA 312
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; TYPE: DNA
; ORGANISM: Chlamydia pneumoniae
US-09-438-185A-1
                CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-6
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 1817
LENGTH: 434
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US-09-438-185A-1/c
                                                                                                                                                                          TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins Patent No. 6783961
FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
                                                                                                                                                                                                                                                                                                                Sequence 1817, Application US/09513999C Patent No. 6783961
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Best Local Similarity 54.7
Matches 152; Conservative
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PRIOR FILING DATE: 1998-11-12
PRIOR APPLICATION NUMBER: US 60/128,606
PRIOR FILING DATE: 1999-04-08
NUMBER OF SEQ ID NOS: 1074
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1
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TYPE: DNA
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CURRENT APPLICATION NUMBER: US/09/438,185A
CURRENT FILING DATE: 2002-03-13
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APPLICANT: Davis, Ronald
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Chlamydia Pneumoniae Genome Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Stephens, APPLICANT: Mitchell,
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Pred. No. 1.4e-13;
0; Mismatches 126;
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; NAME/KEY: UNSURE
; LOCATION: 63
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FEATURE:
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; OTHER INFORMATION: X
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THER INFORMATION:
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OCATION: 293
THER INFORMATION: n=a,
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OCATION: 292
THER INFORMATION: n=a,
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OCATION: 291
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THER INFORMATION: k=g
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CHER INFORMATION: w=a
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                       CAATTTAACTATTGGCGCTCCAGTTATTGAAAGTTTGAGAAAGCA-----CACAAAGGC
                                              GATGCTAGACTCTKGGGCCGATTATCTGCACCTGGACGTAATGGACGGGCATTTTGTTCC
                                                            AAGATTGGCCCGTCCATCCTCAACAGCGACCTGGCCAATTTAKGGGCCGAGTGCCTCCGG
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Pred. No. 2.1e-15;
3; Mismatches 149;
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US-09-902-540-9495
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PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 9495
                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 147; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9495, Application US/09902540 Patent No. 6833447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Stater, Steven C.
APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15849)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/902,540 CURRENT FILING DATE: 2001-07-10
                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 64
TYPE: DNA
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                                TCTGGTTTTACATTTCACGTAGAGAC 301
                                                              GTGCACCTGATGATTGTGGAGCCGGAGCGCTACGTGGAGGC
                                                                                             TGTCACCTTATGGTTACAAATCCTCTTGATTATGTTGAACCCTTGGCAAAAGCTGGTGCT 275
                                                                                                                                ATCACGATTGGCCCCGTGGTGGTGGAGGCCATCAAGCGGGTGGCGACGAAGCCGTTGGAC
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ilarity 55.3%;
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Pred. No. 3e-15;
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Search completed: May 4, 2005, 12:28:18 Job time : 124.536 secs

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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

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Result
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/ Cgn2_6/ptodata/1/pubpna/US09C_PUBCOMB.seq:*
/ Cgn2_6/ptodata/1/pubpna/US09C_PUBCOMB.seq:*
/ Cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
/ Cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
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/ Cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
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jn2_6/ptodata/1/pubpna/USO6_EWE PUB.seq:*
jn2_6/ptodata/1/pubpna/USO6_PUBCOMB.seq:*
jn2_6/ptodata/1/pubpna/USO7_NEW_PUB.seq:*
jn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
jn2_6/ptodata/1/pubpna/USO8_NEW_PUB.seq:*
jn2_6/ptodata/1/pubpna/USO8_NEW_PUB.seq:*
jn2_6/ptodata/1/pubpna/USO8_PUBCOMB.seq:*
jn2_6/ptodata/1/pubpna/USO8_PUBCOMB.seq:*
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US-09-987-899-4761
US-10-424-599-109667
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US-10-021-323-6163
US-09-987-899-4727
US-09-987-899-4725
US-09-987-899-4726
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US-09-987-899-4726
US-10-437-963-45062
US-10-425-114-4370
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                    Sequence 6163, Ap
Sequence 4727, Ap
Sequence 4725, Ap
Sequence 92527, A
Sequence 4726, Ap
Sequence 47062, Ap
Sequence 12139, A
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Sequence 1149, Ap	Sequence 1149, Ap	Sequence 618, App	Sequence 618, App	Sequence 319, App	Sequence 25846, A	Sequence 25186, A	Sequence 12, Appl	Sequence 617, App	Sequence 617, App	Sequence 321, App	Sequence 2058, Ap		Sequence 43914, A	Sequence 4705, Ap	Sequence 7334, Ap				180354	4706	Sequence 4114, Ap	4114,	1137		4723,	4728,	514,	521,	520,	e 513,	Sequence 519, App	Ф	1553

ALIGNMENTS

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CURRENT APPLICATION NUMBER: US/09/987,899
CURRENT FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: US 09/262,979
PRIOR FILING DATE: 1999-03-04
PRIOR APPLICATION NUMBER: US 60/076,712
PRIOR FILING DATE: 1998-03-06
PRIOR FILING DATE: 1998-03-06
NUMBER OF SEQ ID NOS: 7341
SEQ ID NO 4761
LENGTH: 417
TYPE: DNA
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US-09-987-899-4761
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                                                                            Best Local Similarity 100. Matches 388; Conservative
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                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Liu, Jingdong
APPLICANT: Miller, Phillip W.
APPLICANT: O Connell, Keith M.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: With the Carbon Assimilation Pathway
FILE REFERENCE: 16517.258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Cheikh,
APPLICANT: Liu, J
                                                                                                                                                               OTHER INFORMATION: Clone ID: LIB3040-057-Q1-E1-C5
                                                                                                                                                                                                                ORGANISM: Glycine max
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                       1 GGAGAAAGAAAGAAAAGATGGGAATGACACCGAAAATAGCTCCTTCGATGCTCTTCCG 60
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                                                                                               100.0%;
                                                                                               100.0%; Score 388; DB 11; 100.0%; Pred. No. 2.7e-110;
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US-10-424-599-109667
; Sequence 109667, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION;
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APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5322))B
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 108667
                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 837
TYPE: DNA
ORGANISM: Glycine max
FEATURE:
                                                                                                                                                                                                                                                                                                                                              Local Similarity 97.4%;
1es 376; Conservation
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                        GATTATGTTGAGCCCTTGGCAAAAGCTGGTGCTTCTGGTTTTACATTTCATGTAGAGACA
                                           GATTATGTTGAACCCTTGGCAAAAGCTGGTGCTTCTGGTTTTTACATTTCACGTAGAGACA 302
                                                                                    AGTTTGAGAAAGCACACAAAAGGGATATTTGGATTGTCACCTTATGGTTACAAATCCTCTT
                                                                                                                                                                      ATGGACATCATGGATGGGCATTTTGTCCCCAATTTAACTATTGGCGCTCCAGTTATTGAA 182
                                                                                                        AGTTTGAGAAAGCACACAAAGGCATATTTGGATTGTCACCTTTATGGTTACAAATCCTCTT
                                                                                                                                                                                                                                                                           AGAAAGAAAGGAAGGATGGGAGTGACACCGAAAATTGCTCCTTCGATGCTCTCTTCCGAC
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                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                         Score 370; DB 17;
Pred. No. 1.6e-104;
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US-10-424-599-109666 Application US/10424599
Sequence 109666, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (5223) B
CURRENT APPLICATION NUMBER: US/10/424,599
UNMBER OF SEQ ID NO.199666
LENGTH: 845
TYPE: DNA
OPCANTON: Children
TYPE: DNA
OPCANTON: Children
TYPE: DNA
OPCANTON: Children
TYPE: DNA
US-10-021-323-6163
; Sequence 6163, Application US/10021323
; Publication No. US20040123340A1
; GENERAL INFORMATION:
; APPLICANT: Deikman, Jill
; APPLICANT: Feng, Paul C.C.
; APPLICANT: Fincher, Karen L.
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                                                                                                                                                                              GTAGCATTAAAGCCTG 378
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGAAAGAAAGAAAGATGGGAATGACACCGAAAATAGCTCCTTTCGATGCTCTCTTTCCGAC
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97.1%;
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Pred. No. 6.7e-101;
0; Mismatches 11;
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US-09-987-899-4727
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PRIOR FILING DATE: 2000-12-14
NUMBER OF SEQ ID NOS: 17880
SEQ ID NO 6163
LENGTH: 629
                                                                PRIOR APPLICATION NUMBER: US 09/262,979
PRIOR FILING DATE: 1999-03-04
PRIOR APPLICATION NUMBER: US 60/076,712
PRIOR FILING DATE: 1998-03-06
NUMBER OF SEQ ID NOS: 7341
SEQ ID NO 4727
                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4727, Application US/09987899 Publication No. US20040116682A1
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                                                                                                                                                                                          APPLICANT: Cheikh, Nordine
APPLICANT: Liu, Jingdong
APPLICANT: Liu, Jingdong
APPLICANT: Miller, Phillip W.
APPLICANT: O Connell, Keith M.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules
TITLE OF INVENTION: With the Carbon Assimilation Pathway
FILE REFERENCE: 16517.258
CURRENT APPLICATION NUMBER: US/09/987,899
CURRENT FILING DATE: 2001-11-16
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TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(5274)B
CURRENT APPLICATION NUMBER: US/10/021,323
CURRENT FILING DATE: 2001-12-12
CURRENT FILING DATE: 2001-12-12
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ORGANISM: Gossypium hirsutum
TYPE: DNA
ORGANISM: Glycine max
                                            ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTGGAAAGAACTTATCCAAAGAATCAAGTCACATGGCATGATTCCTGGTGTAGCATTAAA 373
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GCACACAAAGGCATATTTGGATTGTCACCTTATGGTTACAAATCCTCTTGATTATGTTGA 253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAGATGGGTGTAACACCAAAAATCGCTCCATCCATGCTTTCATCGGATTTTGCTAACTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GCCTGGGACCCC 385
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GCCTGGAACACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTGGCAAGAACTTATCCAAAAAATTAAGTTGAAAGGCATGAGGCCTGGTGTGGCGTTAAA 451
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81.5%;
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; FEATURE:
; OTHER INFORMATION: Clone ID: 701007767H1
US-09-987-899-4727
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US-09-987-899-4725
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SEQ ID NO 4725
LENGTH: 272
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                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Liu, Jingdong
APPLICANT: Miller, Phillip W.
APPLICANT: O Connell, Keith M.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: With the Carbon Assimilation Pathway
FILE REFERENCE: 16517 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/987,899
CURRENT FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: US 09/262,979
PRIOR FILING DATE: 1999-03-04
PRIOR APPLICATION NUMBER: US 60/076,712
PRIOR FILING DATE: 1998-03-06
                                                                                                                                                                                                                                                                                FEATURE:
NAME/KEY: unsure
LOCATION: (1)...(272)
OTHER INFORMATION: unsure at all n locations
OTHER INFORMATION: Clone ID: 700971857H1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Cheikh, Nordine APPLICANT: Liu, Jingdong
                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Glycine
                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
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hes 257; Conser
                     123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     190 GAAAGCACACAAAGCATATTTGGATTGTCACCTTATTGGTTACAAATCCTCTTGATTATG
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                                                                                                                                                                                                     254;
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ATGGACATCATGGATGGGCATTTTGTCCCCAATTTAACTATTGGCGCTCCAGTTATTGAA 182
                                                                                                                                                                    AGAAAGAAAGAAAGATGGGAATGACACCGAAAATAGCTCCTTCGATGCTCTCTTCCGAC 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTTGGCTTCCGAGGCTCAGCGCATGCTCCACTTCGGCGCCGATTGGCTCCACATGGACA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAGGAAGGATGGGAGTGACACCGAAAATTGCTCCTTCGATGCTCTCTTCCGACTTCGCCA 61
                                                                                          TTCGCCAATTTGGCTTCCGAGGCTCAGCGCATGCTCCACTTCGGCGCCGATTGGCTCCAC 122
                                                                                                                                  AGANAGAAAGGAAGGATGGGAGTGACACCGAAAATTGCTCCTTCGATGCTCTCTTCCGAC 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAAAGCACACAAAGGGATATTTGGATTGTCACCTTATGGTTACAAATCCTCTTGATTATG
                                                         TTCGCCAATTTGGCTTCCGAGGCTCAGCGCATGCTCCACTTCGGCGCCGATTGGCTCCAC 130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTGAGCCCTTGGCAAAAGCTGGTGC 266
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97.0%;
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96.9%;
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Pred. No. 4.8e-68;
                                                                                                                                                                                                                        Score 250.4; DB : Pred. No. 1.8e-67
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RESULT 8
US-09-987-899-4726
VS-09-987-899-4726, Application US/09987899
Publication No. US20040116682A1
GENERAL INFORMATION:
APPLICANT: Cheikh, Nordine
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; FEATURE:
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_184377C.1
US-10-425-115-92527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 7
US-10-425-115-92527
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APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(52322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 92527
LENGTH: 499
TURESTH: 499
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Publication No. US20040214272A1
GENERAL INFORMATION:
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les 295; Conservative
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                                                                                                                                                                   GCCTGGGACCCCCGT 388
                                                                                                                                                                                                                               CTGGAAAGAACTTATCCAAAGAATCAAGTCACATGGCATGATTCCTGGTGTAGCATTAAA 373
                                                                                                                                                                                                                                                                    GGCATTTGGAAAAGCTGGTGCCTCAGGATTCACATTCCATATAGAAGTAGCGAGGGATAA
                                                                                                                                                                                                                                                                                          ACCCTTGGCAAAAGCTGGTGCTTCTGGGTTTTACATTTCACGTAGAGACATCAAAAGATAA
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                                                                                                                                                                                                     CTGGAAAGAGCTCACCCAAAGCATCAAAGCAAAGGGCATGCGGCCTGGTGTATCATTGAA
                                                                                                                                                                                                                                                                                                                                    GCACACAAAGGCATATTTGGACTGCCATCTCATGGTCACAAATCCTTCCGATTATGTAGA 329
                                                                                                                                                                                                                                                                                                                                                         GCACACAAAGGCATATTTGGATTGTCACCTTATGGTTACAAATCCTCTTGATTATGTTGA 253
                                                                                                                                                                                                                                                                                                                                                                                                                          GGATGGGCATTTTGTCCCCAATTTAACTATTGGCGCTCCAGTTATTGAAAGTTTTGAGAAA 193
                                                                                                                                         GCCTGGTACTCCTGT 464
                                                                                                                                                                                                                                                                                                                                                                                                       GGATGGGCACTTCGTTCCTAATTTAACTATTGGAGCTCCAGTGATTGAGAGCTTGAGGAA
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78.7%;
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Pred. No. 2.8e-66;
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; APPLICANT: Liu, Jingdong
; APPLICANT: Miller, Phillip W.
; APPLICANT: Miller, Phillip W.
; APPLICANT: O Connell, Keith M.
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
; TITLE OF INVENTION: With the Carbon Assimilation Pathway
; FILE REFERENCE: 16517.258
; CURRENT APPLICATION NUMBER: US/09/987,899
; CURRENT FILING DATE: 2001-11-16
; PRIOR APPLICATION NUMBER: US 09/262,979
; PRIOR APPLICATION NUMBER: US 09/262,979
; PRIOR FILING DATE: 1999-03-04
; PRIOR FILING DATE: 1999-03-06
; NUMBER: OF SEQ ID NOS: 7341
; SEQ ID NO 4726
; LENGTH: 260
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-437-963-45062

; Sequence 45062, Application US/10437963

; Publication No. US20040123343A1

; GENERAL INFORMATION:
                                                            APPLICANT: Wu, Wei APPLICANT: Boukharov, Andrey A.
APPLICANT: Boukharov, Andrey A.
APPLICANT: Boukharov, Andrey A.
APPLICANT: Boukharov, Andrey A.
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REPERENCE: 38-21(55221)B
FULE REPERENCE: 38-21(55221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NO 45062
SEQ ID NO 45062
LENGTH: 1406
TYPE: DNA
ORGANISM: Oryza sativa
FEATURE:
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; OTHER INFORMATION: Clone ID: 701064495H1
US-09-987-899-4726
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                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
NAME/KEY: unsure
LOCATION: (1)..(1406)
OTHER INFORMATION: un
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Glycine max
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATGGACATCATGGATGGGCATTTTGTCCCCCAATTTAACTATTGGCGCTCCAGTTATTGAA 182
                                                                                                                                                                                                                                                                                                                                                           Cao, You
Wu, Wei
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Pred. No. 8.3e-65;
0; Mismatches 7
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; OTHER INFORMATION:
US-10-437-963-45062
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US-10-767-701-12139
                                                                                                                                                                                                                                                                                                                                APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants and Uses Thereof For Plant Improvement
FILE REFERENCE: 38-21(53535)8
CURRENT APPLICATION NUMBER: US/10/767,701
CURRENT FILING DATE: 2004-01-29
NUMBER OF SEQ ID NOS: 63128
SEQ ID NO 12139
LENGTH: 1026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 12139, Application US/10767701 Publication No. US20040172684A1 GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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TGTCCCCAATTTAACTATTGGCGCTCCAGTTATTGAAAGTTTGAGAAAGCACACAAAGGC
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78.8%;
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76.3%;
                                                                                                                                                                                  Score 225.4; DB 18; Lengtl
Pred. No. 2.3e-59;
0; Mismatches 86; Indels
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US-10-425-114-4370
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APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack B
APPLICANT: Tabaska, Jack B
APPLICANT: Tabaska, Jack B
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Plants and Olecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 4370
LENGTH: 1027
TYPE: DNA
ORGANISM: Zea mays
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   CGT 388
                                                         TATCCAAAGAATCAAGTCACATGGCATGATTCCTGGTGTAGCATTAAAGCCCTGGGACCCC 385
                                                                                                                                                                                                                                    CGTTCCTAACCTGACTATTGGGGCTCCGGTGATCCAGAGCTTGAGGAAACATACCAAAGC
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                                                                                                    GGCTGGCGCTTCTGGATTCACATTCCATATAGAAGTTGCTAGAGACAACTGGCAAGATCT
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76.0%;
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Pred. No. 7.2e-59;
0; Mismatches 87;
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Length 1027;

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RESULT 13
US-10-425-114-28690
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Sequence 28690, Application US/10425114
Publication No. US20040034888A1
GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_7329C.1
US-10-425-115-155395
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US-10-425-115-155395
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(5322)B
FULE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 155395
LENGTH: 1250
TYPE: NAM
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Best Local
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Publication No. US20040214272A1
GENERAL INFORMATION:
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ORGANISM: Zea mays
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276; Conservative
                                                                                                                                                                                                                                                                                                                           CGT 388
                                                                                                                                                                                                                                                                                                                                                                                     TATCCAAAGAATCAAGTCACATGGCATGATTCCTGGTGTAGCATTAAAGCCTGGGACCCC 385
                                                                                                                                                                                                                                                                                                                                                      CATCCAAAGCATTAAATCAAAGGGTATGCGGCCTGGTGTATCATTGAGGCCAGGTACTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGCTGGTGCTTCTGGTTTTTACATTTCACGTAGAGACATCAAAAGATAACTGGAAAAGAACT 325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATATTTGGACTGCCATCTTATGGTCACAAAGCCTTCAGATTACGTAGAACCATTTGGAAA 428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATATTTGGATTGTCACCTTATGGTTACAAATCCTCTTGATTATGTTGAACCCTTGGCAAA 265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CGTTCCTAACCTGACTATTGGGGCTTCCGGTGATCCAGAGCTTGAGGAAACATACCAAAGC
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Pred. No. 8e-59;
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AAAGAAAAGATGGGAATGACACCGAAAATAGCTCCTTCGATGCTCTCTTCCGACTTCGCC

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APPLICANT: Blokeberg, Leonard N.
ITITLE OF INVENTION: Materials and Methods for the
ITITLE OF INVENTION: Modification of Plant Cell Wall Polysaccharides
FILE REFERENCE: 11000.1012c3
CURRENT APPLICATION NUMBER: US/10/393,840
CURRENT FILING DATE: 2003-03-20
PRIOR APPLICATION NUMBER: US 09/636,800
PRIOR APPLICATION NUMBER: US 09/170,862
PRIOR FILING DATE: 1998-10-13
PRIOR FILING DATE: 1998-10-13
PRIOR APPLICATION NUMBER: US 09/170,862
PRIOR FILING DATE: 1998-10-13
PRIOR APPLICATION NUMBER: US 09/148,426
PRIOR FILING DATE: 1999-10-18
PRIOR APPLICATION NUMBER: US 07/148,426
PRIOR FILING DATE: 1999-10-08
                                                                                          US-10-393-840-519
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US-10-393-840-519
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                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 519
LENGTH: 705
                          Query Match
Best Local Similarity
        Matches 275;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 519, Application US/10393840 Publication No. US20030229922A1 GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 28690
LENGTH: 959
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Best Local Similarity
                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 956
                                                                                                       TYPE: DNA
ORGANISM: Pinus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Zea mays
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              426
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    366 GGCTGGCGCTTCTGGATTCACATTCCATATAGAAGTTGCTAGAGACAACTGGCAAGATCT
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      Conservative
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ilarity 76.3%;
Conservative
                     73.5%;
Score 215.6; DB 17;
Pred. No. 2.1e-56;
0; Mismatches 99;
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Pred. No. 1.4e-57;
D; Mismatches 84
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  Indels
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Gaps
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APPLICANT: Bloksberg, Leonard N.
TITLE OF INVENTION: Materials and Methods for the
TITLE OF INVENTION: Modification of Plant Cell Wall Polysaccharides
FILE REFERENCE: 11000.1012c3
CURRENT APPLICATION NUMBER: US/10/393,840
CURRENT FILING DATE: 2003-03-20
PRIOR APPLICATION NUMBER: US 09/636,800
PRIOR FILING DATE: 2000-08-10
PRIOR FILING DATE: 1998-10-13
PRIOR APPLICATION NUMBER: US 09/170,862
PRIOR APPLICATION NUMBER: US 60/148,426
PRIOR APPLICATION NUMBER: US 60/148,426
PRIOR FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: US 60/148,426
PRIOR FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: US 60/148,426
PRIOR FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: US 60/148,426
PRIOR FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: US 60/148,426
PRIOR FILING DATE: 1999-10-08
NUMBER OF SEQ ID NOS: 956
SOFTWARE: FastSEQ for Windows Version 4.0
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US-10-393-840-513
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; LENGTH: 508
; TYPE: DNA
; ORGANISM: Pinus radiata
US-10-393-840-513
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                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                 / Match 55.4%; Score 215; DB 17; Length 5
Local Similarity 73.3%; Pred. No. 2.7e-56;
les 275; Conservative 0; Mismatches 100; Indels
188 GAGAAAGCACAAAAGGCATATTTGGATTGTCACCTTATGGTTACAAATCCTCTTGATTA 247
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                                                                                                                                                                                                                                                                         97
                                                                                                                                                                                                                                                                                                  8 GAAAGAAAAGATGGGAATGACACCGAAAATAGCTCCTTCGATGCTCTCTTCCGACTTCGC
                                                                                                                                                                                                                                                                      GAACACAGAGAAGGGGGTTATCCCTAAAATTGCCCCGTCAATGTTGTCATCAGACTTTGC 156
                                                                                                       CATCATGGATGGGCATTTTGTCCCCAATTTAACTATTGGCGCTCCAGTTATTGAAAGTTT 187
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                                                                                                                                                                                                                                                                                                                                                                                                              Length 508;
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                                                                                                                                                                                                         277 GAGGAAGCATACCCAGGCATTCTTGGATTGTCATCTTATGGTCACAAACCCTCTTGATTA 336
457
                                                                                              AGATAACTGGAAAGAACTTATCCAAAGAATCAAGTCACATGGCATGATTCCTGGTGTAGC 367
                       ATTAMAGCCTGGGAC 382
                                                                                                                                      TGTGGAGCCATTTGCAAAAGCTGGAGCTTCAGGGTTCACTTTTCATGTGGAGGCTGCCAA
AGTGAAACCTGGAAC 471
                                                                 AGACAATTGGCAAGATCTCATCAAAAGAATCAGAAATGCTGGCATGCGGCCTGGAGTGGC
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Search completed: May 4, 2005, 21:11:40 Job time : 443.206 secs

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Minimum
Maximum
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB
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Match Length DB
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2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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                      US-09-583-110-44
US-09-410-5518-1
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422343333434444433	15.0	15.1	15.1	15.1	15.1	15.2	15.2	15.2	15.2	15.2	15.2	15.2	15.3	15.3	15.3	15.3	15.3	
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	US-09-252-991A-13731	US-09-103-840A-2	US-09-902-540-1172	US-09-902-540-3588	US-10-101-464A-286	US-09-899-642A-1	US-09-380-420C-1	US-09-072-967-37	US-09-072-596-37	US-09-056-556-37	US-08-818-111-37	US-08-818-112-37	US-09-635-359B-1	US-09-231-818-1	US-08-510-646B-1	US-08-403-852D-1	US-09-252-991A-5584	

ALIGNMENTS

US-09-583-110-44

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; NUMBER OF SEQ ID NOS: 5322
; SEQ ID NO 44
; LENGTH: 741
; TYPE: DNA
; ORGANISM: Streptococcus pneumoniae
US-09-583-110-44
                                                                                                                                                                                                                                                               RESULT 2
US-08-440-856A-9
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                                                               Sequence 9, Application US/08440856A
PACENT NO. 5750873
GENERAL INFORMATION:
APPLICANT: DELLAPORTA, STEPHEN L.
TITLE OF INVENTION: MATERIALS AND METHODS FOR PRODUCING
TITLE OF INVENTION: MATERIALS AND METHODS FOR PRODUCING
TITLE OF INVENTION: PLANTS WITH SINGLE-SEX FLOWERS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 67; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 44, Application US/09583110 Patent No. 6699703 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Lynn Doucette-Stamm et al.
TITLE OF INVENTION: Nucleic Acid and Amino Acid Sèquences Relating (
TITLE OF INVENTION: Nucleic Acid and Amino Acid Sèquences Relating (
TITLE OF INVENTION: Phenmoniae for Diagnostics and Therapeutics FILE REFERENCE: PATH00-07A
CURRENT FILING DATE: PATH00-07A
CURRENT FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/107,433
PRIOR FILING DATE: 1998-06-30
PRIOR APPLICATION NUMBER: US 60/085,131
PRIOR PILING DATE: 1998-05-12
PRIOR FILING DATE: 1998-05-12
PRIOR FILING DATE: 1998-05-12
PRIOR FILING DATE: 1997-07-02
PRIOR FILING DATE: 1997-07-02
STREET:
CITY: V
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                                                                                                                                                                                                                                                                                                                                                                                                                 231 CTCGGCCTAGGCACCGGCTCCACCGCCAAGCATGCCGTCGA 271
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Conservative
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Pred. No. 0.0077;
0; Mismatches 3
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SEQ ID NO 1
LENGTH: 7
                                                          APPLICANT: WU, KAI
TITLE OF INVENTION: POLYKETIDE SYNTHASE ENZYMES AND RECOMBINANT DNA
TITLE OF INVENTION: CONSTRUCTS THEREFOR
FILE REFERENCE: 30062-20028.00
CURRENT APPLICATION NUMBER: US/09/410,551B
CURRENT ETLING DATE: 1999-10-01
PRIOR APPLICATION NUMBER: US 60/139,650
PRIOR FILING DATE: 1999-08-17
PRIOR APPLICATION NUMBER: US 60/123,810
PRIOR FILING DATE: 1999-03-11
PRIOR FILING DATE: 1999-03-11
PRIOR FILING DATE: 1999-03-10
PRIOR FILING DATE: 1999-03-10
PRIOR FILING DATE: 1999-03-10
PRIOR FILING DATE: 1999-03-10
PRIOR FILING DATE: 1998-10-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1, Application US/09410551B Patent No. 6503737
                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 86; Conserv
                                                                                                                                                                                                                                                                                                                                                            APPLICANT: KOSAN BIOSCIENCES, Inc.
APPLICANT: REEVES, CHRISTOPHER
APPLICANT: CHU, DANIEL
APPLICANT: KHOSLA, CHAITAN
                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: 706141
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/
FILING DATE: 15-MAY-1995
CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: MILLMAN, ROBERT A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 36,217
REFERENCE/DOCKET NUMBER: 05-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 887-1517
TELEFAX: (202) 887-0763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
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                                      FastSEQ for Windows Version 4.0
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CHU, DANIEL
KHOSLA, CHAITAN
SANTI, DANIEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AGTCCGGCATGGTCCTCGGCCTAGGCACCGGCTCCACCGCCAAGCATGCCGTCGACCG 274
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48.3%; Pred. No. 0.01;
ative 13; Mismatches
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US-09-940-316B-1/c
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                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 77536
                                                                                                                                                                                                                                                                Query Match
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: WU, KAI
TITLE OF INVENTION: POLYKETIDES ENCODING THE FK-520 POLYKETIDE SYNTHI
TITLE OF INVENTION: GENE CLUSTER
FILE REFERENCE: 30062-20026.11
CURRENT APPLICATION NUMBER: US/09/940,316B
CURRENT FILING DATE: 2001-08-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 1998-10-02
NUMBER OF SEQ ID NOS: 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/102,748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 09/410,551
PRIOR FILING DATE: 1999-10-01
PRIOR APPLICATION NUMBER: US 60/139,650
PRIOR FILING DATE: 1999-06-17
PRIOR FILING DATE: 1999-03-11
PRIOR FILING DATE: 1999-03-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Streptomyces hygroscopicus
                                                                                                                                                                                                                                                                                                                      NAME/KEY: CDS
LOCATION: (52)
                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
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LOCATION: (52275)...(71465)
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46231 CACCTCGACCAGGTGCGCCGCACCGGCACCCCACCGTCGTCGTCGTCGGGCGCCCG 46173
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les 95; Conserv
                                                                               46291
                                                                                                                                               46351 TACCTCGCCGCCGCCGCCGCCGCCGTCGCCGTCCCGCTCAACCCGGACGCCGCCGAC
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                           216 GAGTCCGGCATGGTCCTCGGCCTAGGCACCGCCCCAAGCATGCCGTCGACCG 274
                                                                                                          156 GTCATCCTCACCCAAGACGATTTGAAGAAAATCGCCGCCTACAAGGCCGTCGAGTACGTG 215
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95; Conser
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                                                                       GCCGAACTCGCCACTTCCTGACGGACTCCGGCGCGACCGTGCTGGTCACCGACGAAACC 46232
                                                                                                                                                                                                                                                                                                                    (52275) . . . (71465)
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CHU, DANIEL
KHOSLA, CHAITAN
SANTI, DANIEL
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53.1%;
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Pred. No. 0.085;
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; LENGTH: 4411529
TYPE: DNA
; ORGANISM: Mycobacterium tuberculosis
; OTHER INFORMATION: H37Rv
US-09-103-840A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-103-840A-1
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Best Local
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                                                                                                                       SOFTWARE: PatentIn
SEQ ID NO 1
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APPLICANT: WHITE, Owen R.
APPLICANT: VENTER, Claire M.
APPLICANT: VENTER, John C.
TITLE OF INVENTION: DAS SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM TITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                    APPLICANT: WHITE, Owen R.
APPLICANT: FRASER, Claire M.
APPLICANT: VENTER, John C.
TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM
TITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-20007.00
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                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: FLEISCHMAN, Robert D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: CDC 1551
OTHER INFORMATION: "n" bases
OTHER INFORMATION: represent
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                                                                                                                                                Ver. 2.1
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Pred. No. 0.24;
0; Mismatches 94;
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Query Match

16.3%;

Score 44.6;

DB 3;

Length 4411529;

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120 ATGGACGCCGGCCTCCTCCACCCCTCCTCCCCCTCATCCTCACCCAAGACGATTTG 179

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US-09-133-321-1
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                                                                 ; LOCATION:
US-09-133-321-1
Query Match
Best Local S
Matches 73
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Patent No. 6255558
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APPLICANT: Haseloff, Ph.
APPLICANT: Hodge, Sarah
TITLE OF INVENTION: IMP
                                                                                                                                                                                                                           TELEX: N/A
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTAL:
ZIP: 22201-4714
ZIP: 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy compatible
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 14-FEB-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9603069.7
FILING DATE: 14-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 701 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 1:
CORRESPONDENCE ADDRESS:
                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
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                                                                                                       NAME/KEY:
                                                                                                                                       TOPOLOGY:
                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                              NAME: Mitchard, Leonard C.
REGISTRATION NUMBER: 29,009
REFERENCE/DOCKET NUMBER: 604-452
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/133,321
FILING DATE: 12-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
                                                                                                                                                                                                                                                           TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Arlington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                 NAME: Mitchard,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        198 AAGGCCGTCGAGTACGTGGAGTCCGGCCATGGTCCTCGGCCTAGGCACCGGCTCCACCGCC
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l Similarity
73; Conserv
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1100 No. 6255558th Glebe Road,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Haseloff, Phillip J.
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              15.7%;
59.3%;
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0; Mismatches
Score 43; DB 3; Length 701; Pred. No. 0.065; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Version #1.30
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RESULT 9
US-09-252-991A-13860
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US-09-252-991A-13584/c
; Sequence 13584, Application US/09252991A
; Patent No. 6551795
Sequence 13860, Application US/09252991A

Patent No. 6551795

GENERAL INFORMATION:

APPLICANT: Marc J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS

TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

CURRENT APPLICATION NUMBER: US/09/252,991A

CURRENT APPLICATION NUMBER: US/09/252,991A

CURRENT FILING DATE: 1998-02-18

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR FILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142

SEQ ID NO 13860
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PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 13584
LENGTH: 753
TYPE: DNA
ORGANISM: Pseudomonas aeruginosa
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Best Local Similarity
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: ARUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
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                                                                                                                                                                                                                                                                                                                                                                                                                GCATGCCGTCGACCG 274
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1998-02-18
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51.3%; Pred. No. 0.067;
tive 0; Mismatches
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0.067;
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US-09-103-840A-1/c
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; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-13860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: FLEISCHMAN, Robert D.
APPLICANT: WHITE, Owen R.
APPLICANT: WHITE, Owen R.
APPLICANT: VENTER, John C.
TITLE OF INVENTION: DIAM SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM TITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT APPLICATION 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: PATENTIN PORT. 2.1
                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO
                                                                                                                                                                                                                                                                                                                                 Query Match
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Patent No. 6294328
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Best Local
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927320 GÁACÁGCAGGCCGGC
                                                                 927380 GCCAGCGCCGAGGTCGGTTCGGCCGGAGCCGCCGGCCCGCCAGCGCCGGAGCC
                                                                                                                                     927440 GGCCCCGCCGGCCGTTACCGCCGTTGCCGATCAGCCCGGCCTTGCCGCCGGCTCC
                                                                                                                                                                                                             927500 GCCGTTACCGAGCCCGAAGGCGGCTCCACCGGCCGCCGCCGCCGCCGCCGCCGCCGCC
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Local Similarity 51.3%;
les 100; Conservative
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                                                                                                198 AAGGCCGTCGAGTACGTGGAGTCCGGCATGGTCCTCGGCCTAGGCACCGGCTCCACCGCC 257
                                                                                                                                                                    138 CACCCCTCCCCCCCCCCCCCCCCCACCCCAAGACGATTTGAAGAAAATCGCCGCCTAC
                                258 AAGCATGCCGTCGAC 272
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51.3%; Pred. No. 0.067;
ative 0; Mismatches
927306
                                                                                                                                                                                                                                                                                      Score 43; DB 3; Length 4411529;
Pred. No. 0.63;
0; Mismatches 95; Indels 0;
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RESULT 11 US-09-902-540-8186 ; Sequence 8186, Application US/09902540 ; Patent No. 6833447

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US-09-902-540-842
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                                                                                                                                                                                                                                               ; TYPE: DNA
; ORGANISM: Myxococcus xanthus
US-09-902-540-842
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APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15849) B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 8186
                                                                                                                                                                                                                                                                                                            SEQ ID NO 842
LENGTH: 6301
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                               APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
ETITE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses
FILE REPERENCE: 38-10 (15849)B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
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                                                                                                                                                                                    Local Similarity
5041
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                                         136
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                                                                                                                                                                 101;
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TGCACGAAGCCATCCGCACCGGCCGCGCGCGCCACGAGGAGCGGCGGCTGCTCAAACGCG 5100
                                       TCCACCCCTCCTCCCCCCCCCCCCCCCCCCCCCCCCAGACGATTTGAAGAAAATCGCCGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCATCGTTGCCGTCTCCC 1102
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                                                                                                                                                            Score 42.8; DB 4;
Pred. No. 0.13;
0; Mismatches 97;
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US-08-440-856A-2
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                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                 Query Match
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MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIAN Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/440,856A
FILING DATE: 15-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (200
TELEFAX: 706141
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (202) 887-0763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: MILLMAN, ROBERT A.
REGISTRATION NUMBER: 36,217
REFERENCE/DOCKET NUMBER: 05
                                                                                                                                                                                                                                                                               Local
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731
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                              CTCCACCGCCAAGCATGCCGTCGACCG
                                                               CGCCGCCTGCGAGCTCGGCGCCCACGGCATCCGCGTCAACTGCATCTCCCCCTTCGGCGT 730
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CGCCACCCCGATGCTCATCAACGCCTG
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MATERIALS AND METHODS FOR PRODUCING PLANTS WITH SINGLE-SEX FLOWERS
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RESULT 14 US-09-252-991A-6615/c

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Sequence 6683, Application US/09252991A

Patent No. 6551795

GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: APRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 107196,136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 6683
LENGTH: 1029
TYPE: DNA
ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-6683
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US-09-252-991A-6683
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; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-6615
                                                                                  Matches
                                                                                                 Query Match
Best Local Similarity
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 107196.136

CURRENT APPLICATION NUMBER: US/09/252,991A

CURRENT FILING DATE: 1999-02-18

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR FILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142

TYPE-CAMA

LENGTH: 813

TYPE-CAMA
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18 ACCACACATTCAATTTTGAACCTCTGGGACTGGCTAGCTTCAACCTTTAACATTAACATG
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                                                                    Score 42; DB 4; Length 1029; Pred. No. 0.13; 0; Mismatches 130; Indels
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Search completed: May 4, 2005, 12:28:35
Job time : 97.1777 secs

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Published Applications NA:*

1: /cgn2_6/ptodata/1/pubpna/US07 PUBCOMB.seq:*

2: /cgn2_6/ptodata/1/pubpna/US06 NEW PUB.seq:*

3: /cgn2_6/ptodata/1/pubpna/US06 NEW PUB.seq:*

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21: /cgn2_6/ptodata/1/pubpna/US10F_PUBCOMB.seq:*

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Copyright (c) 1993 - 2005 Compugen Ltd.
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Sequence Sequence	7-899-4779 Sequence 4779, 7-899-4780 Sequence 4780,	Sequence Sequence	Sequenci	Sequence	Sequence	Description

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45			ö	•	46.2	46.4	46.6	48	48.2	æ	49.6	50.2	50.2	50.2	50.4	54.6	55.2	59.4	•	•	61	62.4	62.4	۲۷.			4	69.4	74.4	81.4	81.4	86	98
16.4	16.7		16.9	16.9	16.9	16.9	17.0	17.5	17.6	17.6	18.1	18.3	18.3	18.3	18.4	19.9	20.1	21.7	21.7	21.7	22.3	٧.	22.8		٧.	ω	23.6	5	.7	29.7	29.7	31.4	31.4
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18	18	19	19	18	18	18	18	18	15	15	9	18	17	17	17	18	18	18	18	1	18	18	H	18	18	17	18	18	17	11	9	17	18
US-10-297-465A-1	10-739-930-4	-10-	-10-363-	-10-363	US-10-363-345A-31407	US-10-437-963-85516	US-10-437-963-96984	ď		US-10-156-761-719	US-09-734-569-113	US-10-425-115-107438	US-10-425-114-35167	US-10-425-114-6710	US-10-424-599-84599	-10	US-10-767-701-152	US-10-739-930-2973	US-10-739-930-2974	-899	-115-	-10	US-09-987-899-4766	-10-739-930-	-10-425-115-	-10	-10-437-963-	US-10-767-701-12123	US-10-424-599-135774	US-09-938-842A-656	US-09-938-842A-656	US-10-424-599-132085	US-10-425-115-133620
Sequence 1, Appli		e 31408,		31408,	31407,	Sequence 85516, A	Sequence 96984, A	402	æ	e 719	113,		35167,		Sequence 84599, A	97674	152, 1	æ	2974,	4765,	13818	39573,	4766,		Sequence 138187,	Sequence 36084, A		1212	1357	Sequence 656, App	Sequence 656, App	Sequence 132085,	Sequence 133620,

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                                             ; FEATURE:
; OTHER INFORMATION: Clone ID: 700745725H1
US-09-987-899-4776
                  Query Match 100.0%; Score 274; DB 11; Best Local Similarity 100.0%; Pred. No. 6e-75; Matches 274; Conservative 0; Mismatches 0;
CTTACATTCCTTTCTCCACCACACATTCAATTTTGAACCTCTGGGACTGGCTAGCTTCAA 60
                               Length
                  Indels
                                274;
                  0
                  Gaps
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RESULT 3
US-09-987-899-4775
; Sequence 4775, Application US/09987899
; Publication No. US20040116682A1
; GENERAL INFORMATION:
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; OTHER INFORMATION: Clone ID: LIB3167-015-A9_FLI
US-10-425-114-19993
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APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 1993
LENGTH: 1422
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Publication No. US20040034888A1
GENERAL INFORMATION:
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ORGANISM: Glycine max
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Local Similarity 97.8%;
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                                                                                                                                      GCACCGCTCCACCGCCAAGCATGCCGTCGACCG 495
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Pred. No. 8.8e-72;
0; Mismatches 6
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APPLICANT: Cheikh, Nordine
APPLICANT: Liu, Jingdong
APPLICANT: Miller, Phillip W.
APPLICANT: Miller, Phillip W.
APPLICANT: O Connell, Reith M.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: With the Carbon Assimilation Pathway
FILE REFERENCE: 16517.258
CURRENT APPLICATION NUMBER: US/09/987,899
CURRENT FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: US 09/262,979
PRIOR FILING DATE: 1999-03-04
PRIOR FILING DATE: 1999-03-06
NUMBER OF SEQ ID NOS: 7341
SEQ ID NO 4775
LENGTH: 263
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US-10-424-599-32036
; OTHER INFORMATION: Clone ID: PAT_MRT3847_128934C.1
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                                                                                                                                                                         APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Covalic David K
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 32036
LENGTH: 1839
TYPE: DNA
TYPE: DNA
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Publication No. US20040031072A1
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Best Local :
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                                                           NAME/KEY: unsure
LOCATION: (1)..(1839)
OTHER INFORMATION: unsure at all n locations
                                                    FEATURE:
                                                                                                                                    FEATURE:
                                                                                                                                                      ORGANISM: Glycine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Glycine max
FEATURE:
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RESULT 6
US-10-424-599-32035
; Sequence 32035, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
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US-10-739-930-3349
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US-10-739-930-3349
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Publication No. US20040216190A1
GENERAL INFORMATION:
APPLICANT: Kovalic, David K.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH
TITLE OF INVENTION: PLANTS AND USES THEREOF FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53377)B
CURRENT APPLICATION NUMBER: US/10/739,930
CURRENT APPLICATION NUMBER: US/10/739,930
CURRENT FILING DATE: 2003-12-18
NUMBER OF SEQ ID NOS: 11088
SEQ ID NO 3349
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Best Local Similarity
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ORGANISM: Glycine max
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENGTH: 1281
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                                                                                                                                   GCCTAGGCACCGGCTCCACCGCCAAGCATGCCGTCGACCG 274
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Pred. No. 4.5e-61;
0; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                        Score 189.6; DB 18; Length 1281; Pred. No. 1.4e-48; 0; Mismatches 9; Indels 3;
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US-10-424-599-32035
                                                                                                                                                                                                                                                APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILLING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 90924
LENGTH: 244
TYPE: DNA
ORGANISM: Glycine max
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APPLICANT: Cao Yongwei
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 32035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 94.5
Matches 208; Conservative
                                                                             Matches
                                                                                             Query Match
Best Local
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
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                                                                                                                                                                                                  NAME/KEY: unsure
LOCATION: (1)..(244)
OTHER INFORMATION: unsure at all n locations
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
                                                                                                                                                                                                FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         305 CACCCATTAACATTAACATGGCCATTCCCTACCCCCATTTCATCGCCACCGAGAAAGCCG
                                                                           al Similarity
197; Conser
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                    CACCACACATTCAATTTTGAACCTCTGGGACTGGCTAGCTTCAACCTTTAACATTAACAT 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCCTCGGCACCGGCTCCACTGCCAAGCACGCCGTCGACCG 524
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CACCACACATCATGGACAAACCTCGGGGCGNGGCTCGCTTTAAACTCTAACATTAACAT
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                                                                                             58.5%;
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                                                                           Score 160.2; DB 17; Length 244;
Pred. No. 1.3e-39;
0; Mismatches 44; Indels 1;
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Pred. No. 1.5e-48;
D; Mismatches 9
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Sequence 4779, Application US/09987899

Publication No. US20040116682A1

GENERAL INFORMATION:

APPLICANT: Chekkh, Nordine

APPLICANT: Liu, Jingdong

APPLICANT: Miller, Phillip W.

APPLICANT: O Connell, Keith M.

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated

TITLE OF INVENTION: With the Carbon Assimilation Pathway

FILE REFERENCE: 16517.258

CURRENT APPLICATION NUMBER: US/09/987,899

CURRENT FILING DATE: 2001-11-16

PRIOR APPLICATION NUMBER: US 09/262,979

PRIOR FILING DATE: 1999-03-04

PRIOR FILING DATE: 1998-03-06

NUMBER OF SEQ ID NOS: 7341

SEQ ID NO 4779

LENGTH: 291

TYPE: DNA

ORGANISM: Glycine max

FEATURE:
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                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                               Sequence 4780, Application US/09987899 Publication No. US20040116682A1
           APPLICANT: Cheikh, Nordine
APPLICANT: Liu, Jingdong
APPLICANT: Miller, Phillip W.
APPLICANT: O Connell, Keith M.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules
FILE REFERENCE: 16517.258
                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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         CURRENT
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LOCATION: (1)...(291)
OTHER INFORMATION: unsure
OTHER INFORMATION: Clone I
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APPLICATION NUMBER: US/09/987,899
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97.0%;
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ID:
                                                                                                                                                                                                                                                                                                                                                                                                                ; Score 98; DB;
; Pred. No. 2.7e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           all n locations
700562390H1
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2.7e-20;
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PRIOR APPLICATION NUMBER: US 09/262,979
PRIOR ETLING DATE: 1999-03-06
PRIOR APPLICATION NUMBER: US 60/076,712
PRIOR FILING DATE: 1998-03-06
NUMBER OF SEQ ID NOS: 7341
SEQ ID NO 4780
LENGTH: 265
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US-10-739-930-3800
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APPLICANT: KOVALIC, DAVIG K.

APPLICANT: KOVALIC, DAVIG K.

TITLE OF INVENTION: NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH

TITLE OF INVENTION: PLANTS AND USES THEREOF FOR PLANT IMPROVEMENT

FILE REFERENCE: 38-21(53377)B

CURRENT APPLICATION NUMBER: US(10/739,930

CURRENT FILING DATE: 2003-12-18

NUMBER OF SEQ ID NOS: 11088

SEQ ID NO 3800

LENGTH: 1027
                                                                                                                                                                                                                                Matches 106;
                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3800, Application US/10739930 Publication No. US20040216190A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Glycine max
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
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                                                                                                                                                         137
                                                                            197
                                                                                                                   191
                                                                                                                                                                         131 CCTCCTCCACCCCCCCCCCCCCCCCCCCCCCCCCAACACCCAAGACGATTTGAAGAAAAATCGC 190
257 CACTGCTGCCTTCGTCGTC 275
                                    251 CACCGCCAAGCATGCCGTC 269
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                                                                      CGCCGACAAGGCCGTGGAGTCCGTCAAGAGCGGCATGGTCCTCGGCCTAGGCACCGGCTC
                                                                                                        CGCCTACAAGGCCGTCGAGTACGTGGAGTCCCGGCATGGTCCTCGGCCTAGGCACCGGCTC 250
                                                                                                                                               CCACTCCCTCCCCATCCGCGCCATCACCCTCACCCAGGACGACCTCAAGAGACACTCGC 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCTCGGCCTAGGCACCGGCTCCACCGCCAAGCATGCCGTCGACCG 274
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                                                                                                                                                                                                                              0; Mismatches
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                                                                                                                                                                                                                                              Score 86.2; DB 18; Pred. No. 1.7e-16;
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Pred. No. 5.4e-19;
0; Mismatches 7;
                                                                                                                                                                                                                              33;
                                                                                                                                                                                                                                                              Length 1027;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 265;
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Sequence 4781, Application US/09987899; Publication No. US20040116682A1; GENERAL INFORMATION:

APPLICANT: Cheikh, Nordine APPLICANT: Liu, Jingdong APPLICANT: Miller, Phillip W. APPLICANT: O Connell, Keith M. US-09-987-899-4781

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RESULT 12
US-10-425-115-133620
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                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Clone ID: MRT4577_53350C.1
JS-10-425-115-133620
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                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 133620
LENGTH: 705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 133620, Application US/10425115 Publication No. US20040214272A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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                                                                                                                                                                     Matches
                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/987,899
CURRENT FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: US 09/262,979
PRIOR FILING DATE: 1999-03-04
PRIOR APPLICATION NUMBER: US 60/076,712
PRIOR APPLICATION NUMBER: US 60/076,712
PRIOR FILING DATE: 1998-03-06
NUMBER OF SEQ ID NOS: 7341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated TITLE OF INVENTION: With the Carbon Assimilation Pathway FILE REFERENCE: 16517.258
                                                                                                                                                                                                                                                                                         FEATURE:
NAME/KEY: unsure
LOCATION: (1)..(705)
OTHER INFORMATION: unsure at all n locations
                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 705
TYPE: DNA
ORGANISM: Zea mays
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TYPE: DNA
ORGANISM: Glycine max
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OTHER INFORMATION: Clone ID:
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                                                                                                                       131 CCTCCTCCACCCCTCCTCCCCCCCCCCCCCTCACCCCAAGACGATTTGAAGAAAATCGC
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                                                                                                                                                                                         Similarity
                                     CGCCTACAAGGCCGTCGAGTACGTGGAGTCCGGCATGGTCCTCGGCCTAGGCACCGGCTC 250
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                                                                                CCACTCCCTCCCCGCCATCGCCCATCACCCTCACCCAGGACGACCTCAAGAGACTCGC
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75.4%;
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Pred. No. 1.4e-16;
                                                                                                                                                                                    Score 86; DB 18; Length 705; Pred. No. 1.8e-16;
                                                                                                                                                                   Mismatches
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RESULT 13
US-10-424-599-132085
; Sequence 132085, Application US/10424599
; Publication No. US:20040031072A1
                                                             APPLICANT: Wang, Xun
APPLICANT: Zhu, Tong
APPLICANT: Zhu, Tong
APPLICANT: Zhu, Tong
TITLE OF INVENTION: STRESS-REGULATED GENES OF PLA
TITLE OF INVENTION: SAME, AND METHODS OF USE
FILE REFERENCE: SCRIP1300-3
CURRENT APPLICATION NUMBER: US/09/938,842A
CURRENT FILING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: US 60/227,866
PRIOR APPLICATION NUMBER: US 60/264,647
PRIOR APPLICATION NUMBER: US 60/264,647
PRIOR APPLICATION NUMBER: US 60/300,111
PRIOR FILING DATE: 2001-06-22
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                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 656, Application US/09938842A Patent No. US20020160378A1 GENERAL INFORMATION:
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SEQ ID NO 132085
SEQ ID NO 656
LENGTH: 798
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Harper, Jeff
APPLICANT: Kreps, Joel
APPLICANT: Wang, Xun
                                              NUMBER OF SEQ ID NOS: 5379
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LOCATION: (1)..(1146)
OTHER INFORMATION: unsure at all n locations
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Pred. No. 2.1e-16;
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US-09-938-842A-656

Sequence 656, Application US/09938842A

Publication No. US20040009476A9

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Harper, Jeff
APPLICANT: Wang, Xun
APPLICANT: Zhu, Tong
ITILE OF INVENTION: STRESS-REGULATED GENES OF PLANTS, TRANSGENIC PLANTS CONTAINING
ITILE OF INVENTION: SAME, AND METHODS OF USE
FILE REFERENCE: SCRIP1300-3

CURRENT APPLICATION NUMBER: US/09/938,842A

CURRENT FILING DATE: 2001-08-24

PRIOR APPLICATION NUMBER: US 60/227,866

PRIOR APPLICATION NUMBER: US 60/264,647

PRIOR APPLICATION NUMBER: US 60/300,111

PRIOR FILING DATE: 2001-06-22

NUMBER OF SEQ ID NOS: 5379

SEQ ID NO 656

LENGTH: 798

ORGANISM: Arabidopsis thaliana

US-09-938-842A-656
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; ORGANISM: Arabidopsis thaliana
US-09-938-842A-656
                                                                                                                                                                                                                                                                                                                                                    Query Match 29.7%; Score 81.4; DB 11; Length 798; Best Local Similarity 64.7%; Pred. No. 5.1e-15; Matches 121; Conservative 0; Mismatches 66; Indels 0
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                                                                               AATTCGTCGAGTCCGGCATGGTTCTCGGTCTCGGAACCGGCTCCACCGCCAAACACGCCG 196
                                                                                                                        AGTACGTGGAGTCCGGCATGGTCCTCGGCCTAGGCACCGGCTCCACCGCCAAGCATGCCG 267
TCGACCG 203
                                             TCGACCG 274
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Search completed: May 4, 2005, 21:11:46 Job time : 316.867 secs

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Result
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Maximum Match 100%
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Perfect score:
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      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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length: 2000000000
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29.8
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              3 US-09-103-840A-2

3 US-09-103-840A-1

3 US-09-064-693A-25

3 US-09-064-693A-25

3 US-08-961-527-49

4 US-08-961-96-374

4 US-08-961-98-374

4 US-09-28-724-1

4 US-09-28-724-1

4 US-09-107-433-1992

5 PCT-09-107-433-1992

6 US-09-643-990A-1

1 US-09-643-990A-1

2 US-09-107-532A-644

4 US-09-107-532A-644

4 US-09-107-532A-644
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US-08-590-454-1
US-09-252-991A-3308
US-09-252-991A-33240
US-09-252-991A-3326
US-09-382-106-1
US-09-489-039A-1895
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Sequence 18, Appl
Sequence 25, Appl
Sequence 374, App
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Sequence 1992, App
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Sequence 10, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 644, Appli
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Sequence 3308, Ap
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Sequence 4813, Ap
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Sequence 3, Appli	Sequence 1118, Ap	Sequence 2579, Ap	.Sequence 4, Appli	Sequence 37, Appl	Sequence 214, App	Sequence 17, Appl	Sequence 17, Appl	Sequence 3383, Ap	Sequence 3340, Ap	Sequence 3294, Ap	Sequence 3367, Ap	Sequence 239, App	Sequence 1, Appl	Sequence 35, Appl	Sequence 1706, Ap	Sequence 201, App	Sequence 2082, Ap

ALIGNMENTS

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RESULT 1
US-09-012-030-1
                                                           US-09-012-030-1
                                                                                                                                                                                                                                                                                                          .....IUM TYPE: Diskette, 3.5 inch, 1.
COMPUTER: IBM AT-compatible, 80486
OPERATING SYSTEM: MS-DOS version 6.(
SOFTWARE: Wordberfect version 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/012.030
FILING DATE:
CLASSTETT:
Query Match
Best Local Similarity
Matches 240; Conserv
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                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08
PILING DATE: 22-JAN-19
PILING DATE: 22-JAN-19
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                MOLECULE TYPE:
HYPOTHETICAL: N
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 20036
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Keil & Weinkauf
STREET: 1101 Connecticut A
                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Uwe TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                       NAME/KEY:
LOCATION:
                                                                                                                    ORGANISM:
                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                        STRANDEDNESS:
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60..2289
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Score 170.4; DB 2;
Pred. No. 3e-39;
0; Mismatches 116;
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                                                                                                                               Query Match
Best Local Similarity
                                                                                                                     Matches 240;
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                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/01
FILING DATE: 22-JAN-1996
CLASSIFICATION: 435
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: SCHMIDT, Ralf-Michael,
APPLICANT: Uwe
TITLE OF INVENTION: Transketolase
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb storage
COMPUTER: IBM AT-compatible, 80486 processor
OPERATING SYSTEM: MS-DOS version 6.0
                                                                                                                                                                                                                     ORGANISM: FEATURE:
                                                                                                                                                                                                                                                MOLECULE TYPE: CI
HYPOTHETICAL: NO
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: WordPerfect version 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                        NAME/KEY:
                                                                                                                                                                                                                                                                                                   STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                  LENGTH: 2629 base pairs
TYPE: nucleic acid
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       1996
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CCGGCACCAAGCCTGACATCATTGGGTTGGGCACCGGCTCCGAGCTGGAGATCGCGGGCA
                                                      CTGGCAACAACCTGATGTCATTTTGATTGGTACTGGCTCAGAGTTAGAAATTGCTGTCA
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                                                                                                             Score 170.4; DB 2;
Pred. No. 3e-39;
0; Mismatches 116;
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PRIOR PILLING DATE: 1998-07-27
NUMBER OF SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT PILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US/074,788
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 3308
LENGTH: 2007
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                      Matches 199;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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Patent No. 6551795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 2007
TYPE: DNA
ORGANISM: Pseudomonas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2236
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                                                                                           GGCCATTGGCATCGACAAGTTCGGCGCGAGTGCTGCTGCCGGGACGATCTACAAGGAGTA 310
                                                                                                                                                            GATCAGCATCGAGGCCGGGTCCACTCTCGGCTGGCAGAAGTACGTCGGAGCCCAAGGCAA 250
                                                               CATCATCGGCATGACCAGCTTCGGCGAGTCGGCGCCGGCGCGCTGTTCGAGCACTT
CGGCTTCACCCTGGACAACGTCCTGGCGGTAGCCGAG 925
                                                                                                                              CATCGCCATCGAGGCCGCCCATGCCGACTACTGGTACAAGTACGTCGGTCTCGACGGGCG
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                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                   30.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                 Score 116.2; DB 4;
Pred. No. 1.2e-23;
0; Mismatches 138;
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US-09-252-991A-3240

Sequence 3240, Application US/09252991A Patent No. 6551795

RESULT 4

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797 190

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FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 199-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 3326
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CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEG ID NOS: 33142
SEG ID NO 3240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 5
US-09-252-991A-3326/c
                                                                                                      US-09-252-991A-3326
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                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
APPLICANT: INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3326, Application US/09252991A Patent No. 6551795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: MARC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 2229
TYPE: DNA
ORGANISM: Pseudomonas
                                                                                                                   TYPE: DNA ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                           ENGTH: 2709
                                       Local
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                  Score 116.2; DB 4
Pred. No. 1.4e-23;
0; Mismatches 138
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Pred. No. 1.3e-23;
0; Mismatches 138;
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APPLICANT: Jiang, Xinhe
APPLICANT: McDevitt, Damien
APPLICANT: Won Horn, Stephanie
TITLE OF INVENTION: tktA
FILE REFERENCE: GM10236
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Patent No. 6221631
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CURRENT FILING DATE: 1999-08-24
NUMBER OF SEQ ID NOS: 2
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TYPE: DNA
ORGANISM: Pseudomonas
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                                                                               CATCATCGGCATGACCAGCTTCGGCGAGTCGGCGCCGGCCCCGGCGCTGTTCGAGCACTT 1943
                                                                                                                                                                                      GATCAGCATCGAGGCCGGGTCCACTCTCGGCTGGCAGAAGTACGTCGGAGCCCAAGGCAA 250
                             CGGCATCACCGTGGAGAGCATCATTGCAACTGCCAAG
                                                                                                                   GGCCATTGGCATCGACAAGTTCGGCGCGAGTGCTCCTGCCGGGACGATCTACAAGGAGTA 310
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CGGCTTCACCCTGGACAACGTCCTGGCGGTGGGCGAG 1980
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Pred. No. 3.5e-23;
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; ORGANISM: Klebsiella pneumoniae
US-09-489-039A-1895
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US-09-489-039A-1895
                                                                                                                       CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 4813
LENGTH: 2127
    Best Local Similarity
                           Query Match
                                                                                                                                                                                                                                                                                                                            Patent No. 6610836
GENERAL INFORMATION:
APPLICANT: Gary Bro
                                                                                                                                                                                                                                                                                                                                                                 Sequence 4813, Application US/09489039A Patent No. 6610836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1895, Application US/09489039A
PATENT NO. 6610836
GENERRAL INFORMATION:
APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PNEUMONIAGE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
CURRENT FILE REPERENCE: 2709.2004001
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 1909-01-29
RUMBER OF SEQ ID NOS: 14342
SEQ ID NO 1895
LENGTH: 2085
                                                                                                                                                                                                                                                  APPLICANT: Gary Breton et. al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: 2709.2004001
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Best Local Similarity
                                                                              TYPE: DNA
ORGANISM: Klebsiella
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAGGCAAGATCGTCGGCATGACCGGCTATGGCGAATCCGCCCCGGCAGAGCAGTTGTTCC 2023
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ilarity 56.9%;
Conservative
                                                                                pneumoniae
  26.1%;
55.8%;
Score 100.4; DB 4;
Pred. No. 4.6e-19;
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Pred. No. 9.6e-21;
0; Mismatches 148;
            Length 2127;
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US-08-311-731A-127
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; Sequence 127, Application US/08311731A
; Patent No. 6583266
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                                                                                                                                                                               FILING DATE:
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: GATES, EDWARD R.
REGISTRATION NUMBER: 31,616
REFERENCE/DOCKET NUMBER: C004
TELECOMMUNICATION INFORMATION:
TELEPHONE: 611/720-3500
TELEPHONE: 611/720-2441
INFORMATION FOR SEQ ID NO: 127:
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                             ORIGINAL SOURCE:
                                              HYPOTHETICAL: 1
                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 35881 base pai
                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: SMITH, DOUGLAS
APPLICANT: MAQ, JEN-I
APPLICANT: SMITH, DOUGLAS
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES
TITLE OF INVENTION: RELATING TO MYCOBACTERIUM TUBERCULOSIS AND LAPRAE FOR
TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                ORGANISM:
                                                                                                      STRANDEDNESS:
TOPOLOGY: cir
                                                                                                                                    TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: UZIP: 02210
                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGAGTTCGGCTTCACCGTTGATAACGTGGTCGCTAAAGCGAA 2114
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGAACTCTTTGATGAGCAGTCGGATGAGTACAAGGAGAGCGTCCTCCCTGCCGACGTCAC 184
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                                                                                                  circular
           MYCOBACTERIUM LEPRAE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: BURKE, KEVIN
APPLICANT: MOCKEL, BETTINA
TITLE OF INVENTION: NEW NUCLEOTIDE SEQUENCES WH
FILE REFERENCE: MAS/21123/258100
CURRENT APPLICATION NUMBER: US/09/531,266
CURRENT FILING DATE: 2000-03-20
PRIOR APPLICATION NUMBER: 60/142,915
PRIOR FILING DATE: 1999-07-09
NUMBER OF SEQ ID NOS: 6
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Best Local Similarity 56.2
Matches 186; Conservative
                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 1
                                                                                                                                                                                                                            Matches 185;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: MCCORMACK, ASHLING APPLICANT: STAPELTON, CLIONA APPLICANT: BURKE, KEVIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: DUNICAN,
                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 6995
TYPE: DNA
ORGANISM: Corynebacterium glutamicum
                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: CDS
LOCATION: (2471)..(3550)
OTHER INFORMATION: tal-Gen
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                         GATGAGCAGTCGGATGAGTACAAGGAGAGCGTCCTCCCTGCCGACGTCACAGCGAGGATC 194
                                                                                               GAGCTGAGGAAGGAGGGAAGACGGTCCGCGTCGTCTCCTCGTCTCCTGGGAACTCTTT
                                                                                                                                                CCAGATGTGATCCTCATGGGCTCCGGCTCCGAGGTTCAGCTTGCAGTTAACGCTGCGAAG
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                                                                        GCTCTGGAAGCTGAGGGCGTTGCAGCTCGCGTTGTTTCCGTTCCTTGCATGGATTGGTTC
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                                                                                                                                                                                                                                       24.4%;
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56.2%;
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                                                                                                                                                                                                                      Score 93.8; DB 4;
Pred. No. 5.9e-17;
0; Mismatches 152;
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US-09-103-840A-2/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 183;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: VENTER, John C.
TITLE OF INVENTION: TUBERCULOSIS
TITLE OF INVENTION: TUBERCULOSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: FLEISCHMAN, Robert D. APPLICANT: WHITE, Owen R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: CDC 1551
OTHER INFORMATION: "n" bases
OTHER INFORMATION: represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Mycobacterium tuberculosis FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ENGTH: 4403765
                                                                                                                                        1628139
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                                                                                                                                        GTCGGCGCGGGTGGCCGTCGAAGCCGGCGTCGCCAATGCTGGCACCAGCTGGTTGGCGA
                                                                                                                                                                                                                                                                                                                                                                          AACCGGCACCAAGCCTGACATCATTGGGTTGGGCACCGGCTCCGAGCTGGAGATCGCGGG
                                CAAGGAGTACGGCATCACCGTGGAGAGCATCATTGCAACTGCCAA 346
                                                                                                    CCAAGGCAAGGCCATTGGCATCGACAAGTTCGGCGCGAGTGCTCCTGCCGGGACGATCTA
                                                                                                                                                                       CACAGCGAGGATCAGCATCGAGGCCGGGTCCACTCTCGGCTGGCAGAAGTACGTCGGAGC
                                                                                                                                                                                                           CCTGGAGTGGTTCGAGGCCCAACCCTACGAGTACCGCGACGCGGTGCTGCCCCCGACGGT
                                                                                                                                                                                                                                          CTGGGAACTCTTTGATGAGCAGTCGGATGAGTACAAGGAGAGCGTCCTCCCTGCCGACGT
                                                                                                                                                                                                                                                                               Application US/09103840A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GTCTCCCTTGAGCACTTCGGTGCTTCTGCGGATTACCAGACCCTGTTTGAGAAGTTCGGC
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CCGCGAGTACGGCTTCACTGCCGAGGCCGTGGCCGCCGCCGCCGA 1627975
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                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           at various positions throughout the sequence {\tt a}, {\tt t}, {\tt c} or {\tt g}
                                                                                                                                                                                                                                                                                                                                                                                                                   Score 85.8; DB 3;
Pred. No. 1.6e-13;
0; Mismatches 162;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 4403765
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RESULT 12 US-09-L03-840A-1/c ; Sequence 1, Application US/09103840A ; Patent No. 6294328

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APPLICANT: FLEISCHMAN, ROBERT D.
APPLICANT: WHITE, OWEN R.
APPLICANT: WHITE, OLA R.
APPLICANT: VENTER, Claire M.
APPLICANT: VENTER, JOHN C.
TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN
TITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 1
SEQ ID NO 3
SE
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US-09-064-693A-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18, Application US/09064693A Patent No. 6210937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Ward, Thomas E.
TITLE OF INVENTION: DEVELC
TITLE OF INVENTION: ENGINE
TITLE OF INVENTION: OF A S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches 183;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: W. Gary Goodson
ADDRESSEE: INEEL--Lockheed M.
ADDRESSEE: Technologies Co.
STREET: P.O. Box 1625
CITY: Idaho Falls
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb
COMPUTER: Toshiba Satellite Pro T2150CDS
OPERATING SYSTEM: Windows95
SOFTWARE: Word Perfect 7.0
CURRENT APPLICATION DATA:
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OTHER INFORMATION: H37Rv
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Pred. No. 1.6e-13;
0; Mismatches 162;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Martin Idaho
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                                                                                                                                                                                                                                                                                                        Sequence 25, Application US/09064693A Patent No. 6210937 GENERAL INFORMATION:
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TELEFAX: (208)526-8339
INFORMATION FOR SEQ ID NO: 18
                                                                                                                                                                                                                    APPLICANT: Ward, Tho
ZIP: 83415-3810
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb
COMPUTER: Toshiba Satellite Pro T2150CDS
OPERATING SYSTEM: Windows95
                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: W. Garry Goodson
ADDRESSEE: INEEL--Lockheed
ADDRESSEE: Technologies Co.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: W. Gary Goodson
REGISTRATION UNMBER: 22,387
REFERENCE/DOCKET NUMBER: LI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                  STREET: P.O. Box 1625
CITY: Idaho Falls
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Simi
                                                                                  COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            GCATCACCGTGGAGAGCATCATTGCAACTGCCAA 346
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                                                                                                                                                                                                                                                                                         Thomas E
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                                                                                                                                                                                                                                DEVELOPMENT OF GENETICALLY ENGINEERED BACTERIA FOR PRODUCTION OF A SPECIFIC PLASTICS PRECURSOR
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Pred. No. 5.6e-14;
0; Mismatches 157;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LIT-PI-296
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                                 storage
                                                                                                                                                                                                                                                        PRODUCTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 2143;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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RESULT 15
US-08-961-527-49
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                                                                                                                                                                                                                                                    Sequence 49, Application US/08961527
Patent NO. 6420135
GENERAL INFORMATION:
APPLICANT: Charles Kunsch
TITLE OF INVENTION: Streptococcus
NUMBER OF SEQUENCES: 391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 177;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (208)526-8339
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: W. Gary Goodson
REGISTRATION NUMBER: 22,387
REFERENCE/DOCKET NUMBER: LI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Word Perfect 7.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                          STREET: STREET: CTTY: Rockville
            MEDIUM TYPE: Diskette, 3.50 inch
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version
SOFTWARE: ASCII Text
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: dou
TOPOLOGY: linear
                                                                                                                    ZIP: 20850
                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3948
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3888 TTGACAAGCAGGATGCTTGCTTACCGTGAATCCGTACTGCCGAAAGCGGTTACTGCACGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13
APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCTTCACTGTTGATAACGTTGTTGCGAAAGCAAA 4101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CCATTGGCATCGACAAGTTCGGCGCGAGTGCTCCTGCCGGGACGATCTACAAGGAGTACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GCATCACCGTGGAGAGCATCATTGCAACTGCCAA 346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TCGTCGGTATGACCACCTTCGGTGAATCTGCTCCGGCAGAGCTGCTGTTTGAAGAGTTCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TTGCTGTAGAAGCGGGTATTGCTGACTACTGGTACAAGTATGTTGGCCTGAACGGTGCTA 4007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TCAGCATCGAGGCCGGGTCCACTCTCGGCTGGCAGAAGTACGTCGGAGCCCAAGGCAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TTGATGAGCAGTCGGATGAGTACAAGGAGAGCGTCCTCCCTGCCGACGTCACAGCGAGGA 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAAAACTGACTGCCGAAGGCGTGAAAGCGCGCGTGGTGTCCATGTCGTCTACCGACGCAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGCCGGAACTGATTTTCATCGCTACCGGTTCAGAAGTTGAACTGGCTGTTGCTGCCTACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGCCTGACATCATTGGGTTGGGCACCGGCTCCGAGCTGGAGATCGCGGGCAATGCGGCCG 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: Human Genome Sciences,
9410 Key West Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6641 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (208) 526-9469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   double
                                                                                                                                                                                                                                                          Streptococcus pneumoniae Polynucleotides and Sequences 391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US/09/064,693A
                                                                             3.50 inch, 1.4Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 82.8; DB 3;
Pred. No. 8.7e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LIT-PI-296
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 6641;
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Best Local Similarity 51.9
Matches 182; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (301) 309-851
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 11443 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 36 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Brookes, A. Anders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE:
                                                                                                                                                                                                                   10123 TGCTCAAGATGCTACCTACAAGGAAGACATTTTACCATCTAAGACTCGTCGTCGTGTGGC 10182
                                                                                                                                                                                                                                                                                                                                                                   10003
10303 TACGGTAGAGAATATCGTTGCTCAAGTTAAGTCCCTATAGAAACCAATTAC 10353
                                                                       10183 CATTGAAATGGCAGCGACCCAAAGTTGGTACAAGTATGTTTGGTTTGGATGGCGCGGTCAT
                                                                                                                                                                                                                                                                                           10063 ATTGGTTTTACAAGGTGGTAAAGTACGTGTGGTATCTATGCCCTCAACCGAACTATTTGA 10122
                                                                                                                                                                                                                                                         137
                                                                                                                                                                                 197
                                   317
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                                                                                                                                                                                                                                                                                                                             TGGCATCGACAAGTTCGGCGAGTGCTCCTGCCGGGACGATCTACAAGGAGTACGGCAT 316
                                                                                                                                                                                                                                                                                                                                                                   TGATACTATTATCATTGCTACAGGATCTGAGGTCAATCTAGCTATCAAAGCTGCTAAGGA 10062
                                                                                                                                                                                                                                                                                                                                                                                                     TGACATCATTGGGTTGGGCACCGGCTCCGAGCTGGAGATCGCGGCCAATGCGGCCGACGA 76
                                  CACCGTGGAGAGCATCATTGCAACTGCCAAGAGCTTTTAAGAGCTAACAAC 367
                                                                                                                                                                               CATCGAGGCCGGGTCCACTCTCGGCTGGCAGAAGTACGTCGGAGCCCAAGGCAAGGCCAT 256
                                                                                                                                                                                                                                                      TGAGCAGTCGGATGAGTACAAGGAGAGCGTCCTCCCTGCCGACGTCACAGCGAGGATCAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 80.6; DB 3;
Pred. No. 4.7e-13;
0; Mismatches 169;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 11443;
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0

Search completed: May 4, 2005, 12:28:49 Job time: 126.658 secs

10242

196

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Minimum DB
Maximum DB
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            seq length: 0
seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-300-482-356
385
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5642217 segs, 3043843248 residues
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Published_Applications_NA:*
|: /cgn2_6/ptodata/1/pubpna/US07_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
_6/ptodata/1/pubpna/US11_NEW_PUB.seq:*
_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
_6/ptodata/1/pubpna/US60_PUBCOMB.seq:*
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                                                                                                       ptodata/1/pubpna/US10F_PUBCOMB.seq:*
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/ptodata/1/pubpna/US10C_PUBCOMB.seq:*
/ptodata/1/pubpna/US10D_PUBCOMB.seq:*
/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
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'1/pubpna/US06_PUBCOMB.seq:*
'1/pubpna/US07_NEW_PUB.seq:*
'1/pubpna/PCTUS_PUBCOMB.seq:*
'1/pubpna/US08_NEW_PUB.seq:*
'1/pubpna/US08_PUBCOMB.seq:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07_PUBCOMB.seq:*
I_NEW_PUB.seq:*
06_NEW_PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.

No.	Score	Query Match	Query Match Length DB	DB	ID	Description
1	385	100.0	385	11	ÜS-09-987-899-4513	Sequence 4513, Ap
N	363.2	94.3	1656	17	US-10-425-114-609	Sequence 609, App
W	361	93.8	1420	17	US-10-425-114-19700	Sequence 19700, A
4	359.4	93.4	3311	18	US-10-425-115-157267	Sequence 157267,
IJ	356.8	92.7	1686	17	US-10-425-114-3351	Sequence 3351, Ap
თ	326.8	84.9	423	11	US-09-987-899-4512	Sequence 4512, Ap
7	295.2	76.7	300	H	US-09-987-899-4502	Sequence 4502, Ap
8	292.2	75.9	2107	18	US-10-767-701-14710	Sequence 14710, A
9	265	68.8	273	11	US-09-987-899-4499	Sequence 4499, Ap
10	249.6	64.8	276	11	US-09-987-899-4501	Sequence 4501, Ap
11	247.8	64.4	646	17	US-10-333-184-131	Sequence 131, App

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444 443 5443	4 2 2 2 2 2 2 4 4 2 2 4 4 2 4 4 4 4 4 4	225 225 226 227 228 232 232 232 232 232 2332	113 114 115 116 117
155 155 155 150.8	163 162.8 161.8 161.8 161.4 160.8	209.2 189.4 183.8 183.8 178.8 178.6 171.6 171.6 171.6 171.6	247.8 247.6 246.4 245.2 241.2 241.2 232 232 230.4
40.3 40.3 9.2			64.4 64.3 64.3 60.3 60.3
2226 2226 2226 2227	1172 1239 693 15172 15172 2608 647	242 235 1758 2354 1146 630 978 2168 2168 3914 2594 2594 1172	1122 2636 630 2869 1127 299 630 630 622
9 9 11 11	17 17 17 17 17	111 117 118 118 117 117 117	17 18 18 18 17 17 17
000.		87-899-4505 87-899-4477 60-238-5947 60-238-5947 237-963-88892 24-599-32349 25-114-969 25-114-9339 24-599-96112 24-599-96112 87-894-334	US-10-333-184-361 US-10-437-963-84321 US-10-767-701-10148 US-10-739-930-5298 US-10-333-184-156 US-09-987-899-4504 US-09-910-664-118 US-10-333-184-138 US-10-333-184-139 US-10-333-184-139
Sequence 150, App Sequence 150, App Sequence 283, App Sequence 283, App Sequence 4506, Ap	135 280 168 32, 34, 129	450 4477 5947 8888 3234 3177 969 933 9611 334 450	Sequence 361, App Sequence 84321, A Sequence 10148, A Sequence 156, App Sequence 156, App Sequence 4504, App Sequence 118, App Sequence 118, App Sequence 139, App Sequence 131, App

ALIGNMENTS

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PRIOR FILING DATE: 1999-03-04
; PRIOR APPLICATION NUMBER: US 60/076,712
; PRIOR FILING DATE: 1998-03-06
; NUMBER OF SEQ ID NOS: 7341
; SEQ ID NO 4513
; LENGTH: 385
; TYPE: DNA
; ORGANISM: Zea mays
; PEATURE: OTHER INFORMATION: Clone ID: LIB3078-007-Q1-K1-G3
US-09-987-899-4513
밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-987-899-4513; Sequence 4513; Application US/09987899; Publication No. US20040116682A1; GENERAL INFORMATION:
                                                                                            Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 385; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Liu, Jingdong
APPLICANT: Miller, Phillip W.
APPLICANT: Ocnnell, Keith M.
APPLICANT: O Connell, Keith M.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: With the Carbon Assimilation Pathway
FILE REFERENCE: 16517.258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/987,899
CURRENT FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: US 09/262,979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Cheikh, Nordine
                                 CAACCGGCACCAAGCCTGACATCATTGGGTTGGGCACCGGCTCCGAGCTGGAGATCGCGG 60
                                                                                                  0
                                                                                            Score 385; DB 11;
Pred. No. 1.8e-108;
); Mismatches 0;
                                                                                                                                           Length 385;
                                                                                                  Indels
                                                                                            0,
                                                                                                  Gaps
60
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APPLICANT: Zhou, Yihua

APPLICANT: Kovalic, David K.

APPLICANT: Screen, Steven E

APPLICANT: Tabaska, Jack E

APPLICANT: Tabaska, Jack E

APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

FILE REFERENCE: 38-21(5:331)B

CURRENT APPLICATION NUMBER: US/10/425,114

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 73128

LENGTH: 1656

TYPE: DNn

TYPE: DNn
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                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Zea mays
FEATURE: THEORY
OTHER INFORMATION: Clone ID: 700084322_FLI
US-10-425-114-609
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milarity 96.6%;
Conservative (
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Pred. No. 1.2e-101;
0; Mismatches 13;
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APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 19700
LENGTH: 1420
TYPE: Nam
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; OTHER INFORMATION: Clone ID: LIB3150-102-F10_FLI
US-10-425-114-19700
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Best Local
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TAACAACGGTCTGAGTTTTTTTATT
                                               ACAAGGAGTACGGCATCACCGTGGAGAGCATCATTGCAGCTGCCAAGAGCTTTTAAGAGC
                                                                  ACAAGGAGTACGGCATCACCGTGGAGAGCATCATTGCCAACTGCCAAGAGCTTTTAAGAGC
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96.1%;
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Pred. No. 5.5e-101;
0; Mismatches 15;
1174
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RESULT 4
US-10-425-115-157267
; Sequence 157267, Application US/10425115
; Publication No. US20040214272A1

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                                        APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53313) B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT APPLICATION NUMBER: US/10/425,114
SEQ ID NO 3351
SEQ ID NO 3351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yinua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
FURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 157267
LENGTH: 3311
                                                                                                                                                                                                                                                                            Sequence 3351, Application US/10425114 Publication No. US20040034888A1
                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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   ORGANISM: Zea mays
                                   ENGTH: 1686
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Pred. No. 2e-100;
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                                                                                                                              Query Match
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; OTHER INFORMATION: Clone ID: LIB3060-013-Q1-K1-A12 US-09-987-899-4512
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; OTHER INFORMATION: Clone ID: 700262031_FLI
US-10-425-114-3351
                                                                                                                                                                                                                                      ; NUMBER OF SEQ ID NOS: 7; SEQ ID NO 4512; LENGTH: 423; TYPE: DNA ORGANISM: Zea mays; FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4512, Application US/09987899 Publication No. US20040116682A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Liu, Jingdong
APPLICANT: Miller, Phillip W.
APPLICANT: Miller, Phillip W.
APPLICANT: O Connell, Keith M.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: With the Carbon Assimilation Pathway
FILE REFERENCE: 16517.258
CURRENT APPLICATION NUMBER: US/09/987,899
CURRENT FILING DATE: 2001-11-16
FRIOR FILING DATE: 1399-03-04
PRIOR PILING DATE: 1399-03-04
PRIOR PILING DATE: 1398-03-06
PRIOR FILING DATE: 1398-03-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Cheikh, Nordifne APPLICANT: Liu, Jingdong
                                                                                           Local Similarity nes 337; Conserv
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                           CAACCGGCACCAAGCCTGACATCATTGGGTTGGGCACCGGCTCCGAGCTGGAGATCGCGG
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CGACCGGCAACAAGCCTGACCTCATTGTGATGGGCACCGGCTCCGAGCTGGAGATCGCGG 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TAACAACGGTCTGGAGTTTTTTT 1502
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                                                                                                                 84.9%;
95.2%;
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95.6%;
                                                                                        Score 326.8; DB 11;
Pred. No. 1.5e-90;
0; Mismatches 17;
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                                                                                                                                        Length 423;
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Sequence 4502, Application US/09987899
Publication No. US20040116682A1
GENERAL INFORMATION:
APPLICANT: Chekk, Nordine
APPLICANT: Chekk, Nordine
APPLICANT: Miller, Phillip W.
APPLICANT: O Connell, Keith M.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: With the Carbon Assimilation Pathway
FILE REFERENCE: 16517.258
CURRENT APPLICATION NUMBER: US/09/987,899
CURRENT APPLICATION NUMBER: US/09/987,899
CURRENT FILING DATE: 1999-03-04
PRIOR APPLICATION NUMBER: US 09/262,979
PRIOR APPLICATION NUMBER: US 09/267,712
PRIOR APPLICATION NUMBER: US 60/076,712
PRIOR FILING DATE: 1998-03-06
NUMBER OF CONTAINS NOT THE NOT TO NOT T
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Best Local :
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LENGTH: 300
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                                                                                                                                             GGCCATTGGCATCGACAAGTTCGGCGCGAGTGCTCCTGCCGGGACGATCTACAAGGAGTA 310
                                                                                                                                                                                                                   GATCAGCATCGAGGCCGGGTCCACTCTCGGCTGGCAGAAGTACGTCGGAGCCCCAGGGCAA
                                                                                                                                                                                                                                                               GATCAGCATCGAGGCCGGGTCCACTCTCGGCTGGCAGAAGTACGTCGGAGCCCAAGGCAA 250
                                                                                                                                                                                                                                                                                                                                CTTTGATGAGCAGTCGGATGAGTACAAGGAGAGCGTCCTCCCTGCCGCCGTCACAGCGAG 120
                                CGGCATCACCGTGGAGAGCATCATTGCAACTGCCAAGAGCTTTTAAGAGCTAACAACGT
                                                                                                                                                                                                                                                                                                                                                                         CTTTGATGAGCAGTCGGATGAGTACAAGGAGGGCGTCCTCCCTGCCGACGTCACAGCGAG 190
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99.0%;
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Pred. No. 7.9e-81;
0; Mismatches 3
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RESULT

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US-09-987-899-4499; Sequence 4499, Application US/09987899; Publication No. US20040116682A1
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US-10-767-701-14710
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Publication No. US20040172684A1

GENERAL INFORMATION:

APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof For Plant Improvement
FILE REFERENCE: 38-21(53535)8

CURRENT APPLICATION NUMBER: US/10/767,701

CURRENT FILING DATE: 2004-01-29

NUMBER OF SEQ ID NOS: 63128

SEQ ID NO 14710

LENGTH: 2107
APPLICANT: Cheikh, Nordine
APPLICANT: Liu, Jingdong
APPLICANT: Liu, Jingdong
APPLICANT: Miller, Phillip W.
APPLICANT: Miller, Phillip W.
APPLICANT: O Connell, Keith M.
TITLE OF INVENTION: With the Carbon Assimilation Pathway
FILE REFERENCE: 16517.258
FILE REFERENCE: 16517.258
CURRENT APPLICATION NUMBER: US/09/987,899
CURRENT FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: US 09/262,979
PRIOR FILING DATE: 1999-03-04
PRIOR APPLICATION NUMBER: US 60/076,712
PRIOR APPLICATION NUMBER: US 60/076,712
PRIOR FILING DATE: 1998-03-06
PRIOR FILING DATE: 1998-03-06
PRIOR APPLICATION NUMBER: US 60/076,712
PRIOR FILING DATE: 1998-03-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 327; Conservative
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84.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 292.2; DB 18; Pred. No. 9.3e-80; Indels
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APPLICANT: Liu, Jingdong
APPLICANT: Miller, Phillip W.
APPLICANT: Miller, Phillip W.
APPLICANT: O Connell, Keith M.
ITILE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated
ITILE OF INVENTION: With the Carbon Assimilation Pathway
FILE REFERENCE: 16517.258
CURRENT APPLICATION NUMBER: US/09/987,899
CURRENT FILING DATE: 2001-11-16
FRIOR APPLICATION NUMBER: US 09/262,979
FRIOR APPLICATION NUMBER: US 09/262,979
PRIOR FILING DATE: 1999-03-04
PRIOR FILING DATE: 1999-03-04
PRIOR FILING DATE: 1998-03-06
NUMBER OF SEQ ID NOS: 7341
SEQ ID NO 4501
LENGTH: 276
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US-09-987-899-4501
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; OTHER INFORMATION: Clone ID: 700044755H1
US-09-987-899-4499
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Best Local Similarity
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LENGTH: 273
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TYPE: DNA
                                                                                                                                                                                                                                                                                                                              ORGANISM: Zea mays FEATURE:
                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Clone ID: 700027425H1
                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
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                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTCGTTCGTCTCCTGGGAACTCTTTGATGAGCAGTCGGATGAGTACAAGGAGAGCGTCCT
TTGATGAGCAGTCGGATGAGTACAAGGAGAGCGTCCTCCCTGCCGACGTCACAGCGAGGA 192
                                                                              CGAGCTGAGG-AAGGAGGGGAAGACGGTCCGCGTCCTCCTTCGTCTCCTGGGAACTCT 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGGGACGATCTACAAGGAGTACGGCATCACCGT 322
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                                                                                                                                                                                                        Conservative
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Pred. No. 1.6e-71;
0; Mismatches 5;
                                                                                                                                                                                                    Score 249.6; DB 11; Length Pred. No. 8.7e-67; O; Mismatches 9; Indels
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APPLICANT: Weglarz, Ted
APPLICANT: Blakslee, Beth
APPLICANT: Oriedo, Jeremiah
APPLICANT: Savickas, Philip
APPLICANT: McCrery, David
APPLICANT: McCrery, David
APPLICANT: Miller, Barbara
APPLICANT: Miller, Barbara
APPLICANT: Miller, Barbara
CURRENT MILLER DOW-04681
CURRENT APPLICATION NUMBER: US/10/333,184
CURRENT APPLICATION NUMBER: 5003-10-10
PRIOR APPLICATION NUMBER: 60/219,809
PRIOR FILING DATE: 2003-07-20
NUMBER OF SEQ ID NOS: 571
SOFTWARE: PatentIn version 3.0
SOFTWARE: PatentIn version 3.0
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; TYPE: DNA
; ORGANISM: Oryza japonica
US-10-333-184-131
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US-10-333-184-131
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                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 291;
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                                                                                             CCCAAGGCAAGGCCATTGGCATCGACAAGTTCGGCGCGAGTGCTCCTGCCGGGACGATCT 300
                                                                                                                                                             TCACAGCGAGGATCAGCATCGAGGCCGGGTCCACTCTCGGCTGGCAGAAGTACGTCGGAG 240
                                                                                                                                                                                                                                 TCAGCATCGAGGCCGGGTCCACTCTCGGCTGGCAGAAGTACGTCGGAGCCCAGGGCAAGG 240
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                                                                     GCAAAGGCAAGGCTATTGGCATCGACAAATTCGGTGCAAGTGCTCCTGCTGGAAAGATCT
                                                                                                                                         TTACTGCAAGAGTCAGCCTTGAAGCAGGGTCTACTCTTGGATGGCAGAAGTACGTCGGAA 409
                                                                                                                                                                                                             GCTGGGAGCTTTTCGATGAACAGTCGGCTGAGTACAAGGAGAGTGTTCTCCCCTGAGGCTG
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Shukla, Vipula
Bauer, Teresa
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Skokut, Thomas
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80.2%;
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Pred. No. 3.6e-66;
0; Mismatches 72
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APPLICANT: Ruegger, Max
APPLICANT: Shukla, Vipula
APPLICANT: Shukla, Vipula
APPLICANT: Blaker, Tere
APPLICANT: Blakslee, Beth
APPLICANT: Blakslee, Beth
APPLICANT: Oriedo, Jeremiah
APPLICANT: Oriedo, Jeremiah
APPLICANT: Savickas, Philip
APPLICANT: Miller, Barbara
APPLICANT: Miller, Barbara
APPLICANT: Miller, Barbara
APPLICANT: Miller, Barbara
APPLICANT: Holler, Barbara
APPLICANT: OF INVENTION: Nucleic Acid Compositions Conferring Dwarfism Phenotype
FILE REFERENCE: DOM-04681
CURRENT FILING DATE: 2003-10-10
PRIOR APPLICATION NUMBER: US/10/333,184
CURRENT FILING DATE: 2000-07-20
PRIOR FILING DATE: 2000-07-20
NUMBER OF SEQ ID NOS: 571
COFFMANDER: DESCRIPTION OF SEQ ID NOS: 571
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Synthetic
US-10-333-184-361
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US-10-333-184-361
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.0 SEQ ID NO 361 LENGTH: 1122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                               TAA 363
                                                                                                                                                                                                                                TCACAGCGAGGATCAGCATCGAGGCCGGGTCCACTCTCGGCTGGCAGAAGTACGTCGGAG 240
                                                                                                                                                                                                                                                                                             Similarity
                                                                                        ACAAGGAGTACGGCATCACCGTGGAGAGCATCATTGCAACTGCCAAGAGCTTTTAAGAGC 360
                                                                                                                               GCAAAGGCAAGGCTATTGGCATCGACAAATTCGGTGCAAGTGCTCCTGCTGGAAAGATCT
                                                                                                                                                   CCCAAGGCAAGGCCATTGGCATCGACAAGTTCGGCGCGAGTGCTCCTGCCGGGACGATCT
CAA 848
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Larrinua, באיי
בייד, Thomas
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 247.8; DB 17; Pred. No. 4e-66; 0; Mismatches 72;
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APPLICANT: Boukharov, Andrey A.

APPLICANT: Li, Ping

TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With

FILE REFRENCE: 38-21 (53221) B

CURRENT APPLICATION NUMBER: US/10/437,963

CURRENT FILING DATE: 203-05-14

NUMBER OF SEQ ID NOS: 204966

ENGTH: 2636

TYPE: No.
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US-10-767-701-10148
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US-10-437-963-84321
Sequence 10148, Application US/10767701
Publication No. US20040172684A1
GENERAL INFORMATION:
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof For Plant Improvement
FILE REFERENCE: 38-21(53535)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 84321, Application US/10437963
Publication No. US20040123343A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David R.
APPLICANT: Chou, Yihua
APPLICANT: Chou, Yihua
APPLICANT: Chou, Yihua
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Best Local Similarity 80.2%;
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ORGANISM: Oryza sativa
FEATURE:
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; TYPE: DNA
; ORGANISM: Sorghum bicolor
; FEATURE:
; FEATURE:
; OTHER INFORMATION: Clone ID: SORBI-28MAY03-CLUS61383_1
US-10-767-701-10148
                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Clone ID: TRIAE-23APR03-CLUSTER730_1608
US-10-739-930-5298
                                                                                                                                                                                                                                                                                                                                             Sequence 5298, Application US/10739930
Publication No. US20040216190A1
GENERAL INFORMATION:
APPLICANT: Kovalic, David K.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH
TITLE OF INVENTION: PLANTS AND USES THEREOF FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53377)B
CURRENT APPLICATION UNDER: US/10/739,930
CURRENT FILING DATE: 2003-12-18
NUMBER OF SEQ ID NOS: 11088
SEQ ID NO 5298
LENGTH: 2869
TYPE: DNA
ORGANISM: Triticum aestivum
FEATURE:
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Matches
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CURRENT FILING DATE: 2004-01-29
NUMBER OF SEQ ID NOS: 63128
SEQ ID NO 10148
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Best Local Similarity
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                                                                                                                                                                                                                               Similarity
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CCTGGGAACTCTTTGATGAGCAGTCGGATGAGTACAAGGAGAGCGTCCTCCCTGCCGACG 180
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                                         CGACCGACAACAAGCCTGACTACATCATAATGAGCACTGGTTCTGAGGTAGAGATTGCTG
                                                                                                                                                              CAACCGGCACCAAGCCTGACATCATTGGGTTGGGCACCGGCTCCGAGCTGGAGATCGCGG 60
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80.7%;
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                                                                                                                                                                       ACAAGGAGTACGGCATCACCGTGGAGAGCATCATTGCAACTGCCAAGAGCTTTTAAGAGC 360
                                                                                   TCACCGCAAGGATCAGCATTGAGGCTGGCTCTACCCTCGGATGGCAAAAGTATGTCGGAT 2227
ACAAGGAGTACGGCATCACCGCGGAGAACGTCATCGCCGCAGCCAAGAGCCTGTAAGATC 2347
                                                       CCAAGGGCAAGACCATCGGCATCGACAAGTTCGGCGCCAGCGCTCCCGCCCCGAAGATCT
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Search completed: May 4, 2005, 21:11:47
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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ALIGNMENTS

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US-08-232-463-14
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Patent No.
GENERAL IN
                                                                                          TELEFAX: (703)683-4109
TELEX: 899149
INFORMATION FOR SEQ ID NO: 14
SEQUENCE CHARACTERISTICS:
LENGTH: 7218 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                      PILING DATE:
APPLICATION NUMBER: EP 91 114
PILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORNATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 3047
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/232,463
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version:
CURRENT APPLICATION DATA:
                          IMMEDIATE SOURCE:
CLONE: pTZgpt-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: DORNER, F.
APPLICANT: SCHEIFLINGER, F.
APPLICANT: FALKNER, F. G.
TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                           TELEPHONE: (703)683-4109
                                                                TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Alexandria
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No. 5670367
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1800 Diagonal Road, Suite 500
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Query Match

19.4%;

Score 51.2;

DB 1;

Length 7218;

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PRIOR APPLICATION NUMBER: 60/241.755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR PRIOR DATE: 2000-10-03
PRIOR ETLING DATE: 2000-09-08
PRIOR ETLING DATE: 2000-09-08
PRIOR ETLING DATE: 2000-09-08
PRIOR ETLING DATE: 2000-09-08
VUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 12751
ENGTH: 148794
                                                                                                                                                                                               RESULT 3
US-09-792-616-1
          GENERAL INFORMATION:
APPLICANT: PXE International, Inc.
APPLICANT: University of Hawaii
TITLE OF INVENTION: Mutations in a gene encoding an ABC transporter (MRP6)
TITLE OF INVENTION: Pseudoxanthoma Elasticum
FILE REFERENCE: PXE-001
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US-09-949-016-12751
                                                                                                                                                     Sequence 1, Application US/09792616
Patent No. 6780587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
CURRENT FILING DATE: 2000-04-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Human
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Pred. No. 0.2;
0; Mismatches 69;
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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LENGTH: 197336
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GENERAL INFORMATION:
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Matches
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NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1
LENGTH: 107820
TYPE: DNA
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TITLE OF INVENTION: FOLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES
FILE REFERENCE: CL001307
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LOCATION: (1)...(19733
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NAME | KEY: misc feature

OTHER INFORMATION: "n" can be an A or a T or a G
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                                                                                                                                                                                                                                                            Similarity
                                                                   CTCTCTGCACTCCTCCCTTCCCTTTCCCTTCCCATCCCTTCCCTTCCCTCCCTCCCTCCCTCCC 14638
                                                                                                     ATCACAAAACTAACCTATCCGCTTCCCTCCGATCCATTCACTCGCTCCCTCTTAAAACCT 192
                                                                                                                                                   AGGTGCCCAGGCCTGGAGGCTCCTGGTCTCCTTCCTTCCCTTCCCTTCCCTTCCCTTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              h 13.2%;
Similarity 50.0%;
                                   CCTTACGGATCAAATGCTCCCAA 215
CCTCTCTGCACGATGGCTGGCCA 14661
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                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                             (197336)
                                                                                                                                                                                                                                                          13.0%;
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                                                                                                                                                                                                                                     Score 34.2; DB 4; Length 197336;
Pred. No. 1.7;
0; Mismatches 68; Indels 0;
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                                                                                                                   SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 17272
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NAME/KEY: misc_feature
LOCATION: (1).T.(197337)
OTHER INFORMATION: n = A,T,C or G
-09-949-016-14376
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GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
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Patent No. 6812339
                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES FITLE OF INVENTION: WITH HUMAN DISEASE, METHODS FILE REFERENCE: CL001307
CHEPRENT ARTS CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. 6812339
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/231,498 PRIOR FILING DATE: 2000-09-08 NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
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PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
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                                                  ORGANISM: Human
AME/KEY: misc_feature
                                                                                            ENGTH: 234288
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Pred. No. 1.7;
0; Mismatches 68;
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OF DETECTION AND USES
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RESULT 8
US-09-949-016-12594/c
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                       GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14044
                                                                                                         Sequence 12594, Application US/09949016 Patent No. 6812339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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Best Local Similarity
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/231,498 PRIOR FILING DATE: 2000-09-08
CURRENT APPLICATION NUMBER: US/09/949,016
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Pred. No. 0.59;
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Pred. No. 1
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-
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                                                                                                                                                                                                                                                                       NAME/KNY: misc_feature;
LOCATION: (1)...(69709);
OTHER INFORMATION: n = A,T,C or US-09-949-016-16036
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PRIOR FILING DATE: 2000-10-20
PRIOR PELING DATE: 2000-10-20
PRIOR PELING DATE: 2000-10-03
PRIOR PELING DATE: 2000-10-03
PRIOR PELING DATE: 2000-09-08
PRIOR PELING DATE: 2000-09-08
PRIOR PELING DATE: 2000-09-08
PRIOR PELING DATE: 207012
SOFTWARE: PASTSEQ for Windows Version 4.
ENGTH: 69263
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NAME/KEY: misc_feature
LOCATION: (1)...(69263)
OTHER INFORMATION: n = A,T,C or G
US-09-949-016-12594
                                                                                                                                                   Matches
                                                                                                                                                                                                             Query Match
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TYPE: DNA
ORGANISM: Human
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT FILING DATE: PRIOR APPLICATION NIM
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                                                                                                                                                                              Local Similarity
59885 CTGCTTCCAGTGCTAATAAACACATAAGAGAACCAACCGCTTAACTTTAAATTACCAAAT 59826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              59765 CTTC 59762
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                                                          115 CTTCTTCCATCGCTTTTAATCACAAAACTAACCTATCCGCTTCCCCTCCGATCCATTCACT 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       175
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                                                                                                                                            Conservative
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                                                                                                                                  Score 32.8; DI Pred. No. 3; O; Mismatches
                                                                                                                                            <u>,</u>
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                                                                                                                                                                                               DB 4;
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; ORGANISM: Human
US-09-949-016-17553
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; ORGANISM: Human
US-09-949-016-14132
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
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                                                          NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 17553
LENGTH: 106380
                                                                                                                                                                                                                                                                  GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

FILE REFERENCE: CL001307

CURRENT FILING DATE: 109/09/949,016

CURRENT FILING DATE: 2000-04-14

CURRENT FILING DATE: 2000-04-14
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14132
LENGTH: 122772
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Best Local :
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                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
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Patent No. 681233
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
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Similarity 59.8%;
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Pred. No. 4.1;
0; Mismatches 37;
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Query Match

12.3%;

Score

32.6;

DB 4;

Length 106380;

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US-09-949-016-2722
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US-09-509-712B-31/c
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Local Similarity
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TYPE: DNA
ORGANISM: Rattus norvegicus
FEATURE:
NAME/KEY: misc feature
LOCATION: 1-857
OTHER INFORMATION: n = g, a,
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APPLICANT: Rubin, Donald H.
                                                                                                                                                                                                                             Sequence 2722, Application US/09949016 Patent No. 6812339
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PRIOR FILING DATE: 1998-10-08
PRIOR APPLICATION NUMBER: 60/062,021
PRIOR FILING DATE: 1997-10-10
NUMBER OF SEQ ID NOS: 127
             APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CLOO1307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
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APPLICANT: DuBois, Raymond
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ITLE OF INVENTION: Mammalian Genes Involved in Viral
ITLE OF INVENTION: Infection and Tumor Suppression
FILING DATE: 2000-10-03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  TTTTCCGTTTTGGCNCACTCCAAACNTTTCCCCCCTANTCCCNCCGCACCCGTTCNTGTNC
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                                                                                                                                                                                                                                                                                                                                                             NCCTGGAAATTTTTNCGG 110
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Pred. No. 0.41;
0; Mismatches 73;
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0; Mismatches
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US-09-949-016-2722
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SEQ ID NO 2722
LENGTH: 4082
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14464
LENGTH: 14205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14464, Application US/09949016 Patent No. 6812339
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Best Local (
                                                                                                                                                                                                                   Best Local Similarity Matches 77; Conserv
                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILLING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60
PRIOR FILING DATE: 2000-09-0
NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/231,498 PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/237,768 PRIOR FILING DATE: 2000-10-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE:
                                                                                                                                                                                                                                                                                                               ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity nes 77; Conserv
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2238
                                                                                                                                           2118 TCCCGGGTCCCCTCGCCCACCCTAATCCACTCTCCCTTCCCTTTCCCGGATTCCCTCGCTC
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                                                                                                         86 CCGAGATTCTCGCTTCCTCACCTTCAAACCTTCTTCCATCGCTTTTAATCACAAAACTAA 145
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CCTTTCCCAGAGACCCAGAGCCCCTGACCCCC 2269
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Pred. No. 1.3;
0; Mismatches
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Pred. No. 2.5;
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RESULT 15 US-09-248-796A-7005/c ; Sequence 7005, Application US/09248796A ; Patent No. 6747137

```
GENERAL INFORMATION:

APPLICANT: Keith Weinstock et al

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN

TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 107196.13

CURRENT APPLICATION NUMBER: US/09/248,796A

CURRENT FILING DATE: 1999-02-12

PRIOR APPLICATION NUMBER: US 60/074,725

PRIOR APPLICATION NUMBER: US 60/096,409

PRIOR FILING DATE: 1998-02-13

PRIOR FILING DATE: 1998-08-13

PRIOR FILING DATE: 1998-08-13

NUMBER OF SEQ ID NOS: 28208

SEQ ID NO 7005

LENGTH: 279

TYPE: DNA

ORGANISM: Candida albicans

US-09-248-796A-7005
Search completed: May 4, 2005, 12:28:52 Job time : 80.2515 secs
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                                                                                                                                             142 CTAACCTATCCGCTTCCCGATCCATTCACTCCCTCCTTAAAACCTCCTTAC 198
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Minimum DB seq length: 0
Maximum DB seq length: 200000000
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Maximum Match 100%
Listing first 45 summaries
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1: /cgn2 6/ptodata/1/pubpna/US07 PUBCOMB.seq:*

2: /cgn2 6/ptodata/1/pubpna/PCT NEW PUB.seq:*

3: /cgn2 6/ptodata/1/pubpna/US06 NEW PUB.seq:*

4: /cgn2 6/ptodata/1/pubpna/US06 PUBCOMB.seq:*

5: /cgn2 6/ptodata/1/pubpna/US07 NEW PUB.seq:*

6: /cgn2 6/ptodata/1/pubpna/US08 NEW PUB.seq:*

7: /cgn2 6/ptodata/1/pubpna/US08 NEW PUB.seq:*

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9: /cgn2 6/ptodata/1/pubpna/US08 PUBCOMB.seq:*

9: /cgn2 6/ptodata/1/pubpna/US08 PUBCOMB.seq:*
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264
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Copyright (c) 1993 - 2005 Compugen Ltd.
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/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.

Score

Query Match Length DB

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17	17	17	17	17	17	17	17	17	17	17
US-10-424-599-28474	US-10-424-599-86542	US-10-424-599-41040	US-10-424-599-132643	US-10-424-599-33477	US-10-424-599-69148	US-10-424-599-10015	US-10-424-599-43250	US-10-424-599-43245	US-10-424-599-43247	US-10-424-599-43249
Sequence 28474, A		Sequence 41040, A	Sequence 132643,	Sequence 33477, A	Sequence 69148, A	Sequence 10015, A	Sequence 43250, A	Sequence 43245, A	Sequence 43247, A	Sequence 43249, A

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Sequence 121972,	Sequence 6140, Ap		Sequence 216903,	e 1398,	1397,	1398,	1397		9496,		613,	Sequence 282, App		Sequence 381, App	Sequence 12122, A	76,		Sequence 9, Appli	æ	Sequence 34, Appl	Sequence 2, Appli	66	Sequence 1, Appli		17	396,		3388,	1557		15721,	1453	Sequence 58838, A

ALIGNMENTS

US-10-424-599-43249; Sequence 43249, Application US/10424599; Publication No. US20040031072A1

GENERAL INFORMATION:

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                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 43249
LENGTH: 747
TYPE: DNA
ORGANISM: Glycine max
                                                                                    Matches 263; Conservative
                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                                                       PEATURE:
NAME/KE; unsure
LOCATION: (1)...(747)
OTHER INFORMATION: unsure at all n locations
                                                                                                                                                                                           OTHER INFORMATION: Clone ID: PAT_MRT3847_139052C.1
                                                                                                                                                                                                                    FEATURE:
116 TGACTTGCAGCAATTCCCAATGGCTTCCGTTTCCAAGCTCTCAACGCCAAATCCACTTGC 175
                         2 TGACTTGCAGCAATTCCCAATGGCTTCCGTTTCCAAGCTCTCAACGCCAAATCCACTTGC 61
                                                                                                      95.5%;
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                                                                                Score 252; DB 17; Length 747; Pred. No. 4.8e-72; O; Mismatches O; Indels
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                                                                                    Gaps
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US-10-424-599-43247
US-10-424-599-43247, Application US/10424599
i Publication No. US20040031072A1
i GENERAL INFORMATION:
APPLICANT: LA Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Corologwei
APPLICANT: Chou Yihua
CUTILE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21[5322]B
CURRENT APPLICATION UNMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 43247
LENGTH: 1926
RESULT 3
US-10-424-599-43245
US-10-424-599-43245, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
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Best Local Similarity
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ORGANISM: Glycine max
FEATURE:
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                                                                                                                                                                                   GCTTCATGATCTTTA 264
                                                                                                                                                              GCTTCATGATCTTTA 335
                                                                                                                                                                                                                          CTCCTTGCGGATCAAATGCTCCCAAACAGATGGGAATGGAAGTCCTGCAAAGAGGACAGT 320
                                                                                                                                                                                                                                                        CTCCTTACGGATCAAATGCTCCCAAACAGATGGGAATGGAAGTCCTGCAAAGAGGACA-T
                                                                                                                                                                                                                                                                                             TAATCACAAAACTAACCTATCGGCTTCGCTCGGATCGATTCACTCGCTCCCTCAAAAAC 190
                                                                                                                                                                                                                                                                                                                                                                                                                                     GCAATTCCCAATGGCCTCCATTTTCAAGCTCTCAACGCCATCTCCAGCTGCTTCCTTATC 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GCAATTCCCAATGGCTTCCGTTTCCAAGGTCTCAACGCCAAATCCACTTGCTTCCTTATC 70
                                                                                                                                                                                                                                                                                                                                                                    AGAAGCGCTTCGCCCAGAGATTCTCGCTTCCTCCTTCAATCCTTCTACCAACGCTAT
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92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 215.8; DB:
Pred. No. 4.9e-60;
0; Mismatches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 17; Length 1926;
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; OTHER INFORMATION: Clone ID: PAT_MRT3847_139053C.1 US-10-424-599-43250
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          Matches 154;
                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 43250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 43250, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION:
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APPLICANT: Cao Yongwei
TITLE OF INVENTION: Sey Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 43245
LENGTH: 592
TYPE: DNA
ORGANISM: Glycine max
FEATURE:
                                                                                                                                                                                                                                                                                                         APPLICANT: La ROSA Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
FILE REFERENCE: 38-21(53223)B
CURRENT FILING DATE: 2003-04-28
CURRENT FILING DATE: 2003-04-28
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Best Local Similarity
                                                                                                                                          NAME/KEY: unsure LOCATION: (1)..(676 OTHER INFORMATION:
                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Glycine max
FEATURE:
                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                 LENGTH: 676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: unsure
LOCATION: (1)..(592
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TCTTAAAACCTCCTTTATGAGAAAGAAGGGCAGAGTCCATGGTATGATAATCTCTGCAGA 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TCTTAAAACCTCCTTACGGATCAAATGCTCCCAAACAGATGGGAATGGAAGTCCTGCAAA 241
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          Conservative
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                                                                                                                                        unsure at all n locations
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      Score 145; DB 17;
Pred. No. 5.4e-37;
0; Mismatches 15;
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Pred. No. 5.4e-54;
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                                        Length 676;
      Indels
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Gaps
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TTTCCAAGCTCTCAACGCCAAATCCACTTGCTTCCTTATCAGAAGCGCTTCGCCCCCGAG 90

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RESULT 6
US-10-424-599-69148
; Sequence 69148, Application US/10424599
; Publication No. US20040031072A1
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Sequence 10015, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
JUNEAU PILICANT: Cao Yongwei

TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With

FILE REFERENCE: 38-21(5323)B

CURRENT APPLICATION NUMBER: US/10/424,599

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 285684

LENGTH: 305
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TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 10015
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GCGCCAATTCTCCTTCGAGCGCTACTCCTTACAAATCATACCTATCCGATAACCTGTGAT 122
                                                                                                                                                                                                                                                                                                                                                                                             ATGTAAGTCCTGCACTGATGACA 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CCATTCACTCGCTCCCTTAAAAACCTCCTTACGGATCAAATGCTCCCAAACAGATGGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTCCATCACAATGTTCTCACTTACCAAAACCGCTACGCTCAAAAAATTCAAAATTGGTCT 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CCGCTTCCCTCCGATCCATTCACTCCCTCCTCTTAAAACCTCCTTGCG 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCGCTTCCCTCCGATCCATTCACTCGCTCCCTTAAAAACCTCCTTACG 199
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nilarity 68.5%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 100.6; D
Pred. No. 1.5e-
0; Mismatches
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                                                                                               RESULT 8
US-10-424-599-132643
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CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 33477
LENGTH: 322
                                      Sequence 132643, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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APPLICANT: Kovalic David K
APPLICANT: Zhou yihua
APPLICANT: Zhou yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Glycine max
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LOCATION: (1)..(30:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Glycine max
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                              / Match 36.7%;
Local Similarity 84.0%;
nes 121; Conservative
                                                                                                                                                                              121
                                                                                                                                                                                                                   242
                                                                                                                                                                                                                                                                                              182
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                                                                                                                                                                                                       GAGGAC-ATGCTTCATGATCTTTA 264
                                                                                                                                                                              CAGGACAATACTTCATGATCTTTA 144
                                                                                                                                                                                                                                                                                        TCTTAAAACCTCCTTACGGATCAAATGCTCCCAAACAGATGGGAATGGAAGTCCTGCAAA 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            THNTHTHCAGCTATTCCHANTGGNNTCCNTHTCCANGCHNNNAACHNCAAATCHANTTHN 217
                                                                                                                                                                                                                                                     TCTTAAAACCTCCTTACGAATCAAATACTCCCAAACAGATAAAAGATAAAGTGCTGCAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CATCGCGTTTAATCACTAAACTNANCTA 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CATCGCTTTTAATCACAAAACTAACCTA 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TTCCTTATCAGAAGCGCTTCGCCCCCGAGATTCTCGCCTTCCTCACCTTCAAACCTTCTTC 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TGACTTGCAGCAATTCCCAATGGCTTCCGTTTCCAAGCTCTCAACGCCAAATCCACTTGC 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (305)
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Pred. No. 1.7e-22;
0; Mismatches 41;
                                                                                                                                                                                                                                                                                                                                                                                                              Score 96.8; DB 17; Length Pred. No. 2.6e-21; O; Mismatches 22; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      322;
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; OTHER INFORMATION: Clone ID: PAT_MRT3847_13705C:1
US-10-424-599-41040
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US-10-424-599-41040
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                                                                                                                                                                                                                                                                                                                                     APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Chovalic David K
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NO 41040
SEQ ID NO 41040
TYDE: NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 41040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 41040, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
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Best Local Simi
Matches 148; (
                                                                                                                                                                   Matches 114;
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Best Local Similarity
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APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 133643
LENGTH: 376
                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Glycine max FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                     196 TAC-GGATCAAATGCTCCCAAACAGATGGGAATGGAAGTCCTGCAAAGAGGACA-TGCTT 253
                                                                                                         136 ACAAAACTAACCTATCCGCTTCCCTCCGATCCATTCACTCGCTCCCTTAAAACCTCCT 195
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TGCGGGGTCAAATGCTCCCAATCAGATGGGAATGGGAGTGCTGCAAAGAGAGACAGTGCTT 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATCAAATGCTCCCAAACAGATGGGAATGGAAGTCCTGCA 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACTAACCTATCCGCTTCCCTCCGATCCATTCACTCGCTCCTTAAAAACCTCCTTACGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCAGAAAAAGAAAAAG-GACACCATAATCCCCCCTATGACCAACGTGTGAAAGTACCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CGCCCCCGAGATTCTCGCTTCCTCACCTTCAAACCTTCTTCCATCGCTTTTAATCACAAA 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATGGCTTCCGTTTCCAAGCTCTCAACGCCAAATCCACTTGCTTCCTTATCAGAAGCGCTT 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AACAAATGCTCCCGTACAGATTGGCACGGATGACCTACA 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATGGCCTCCATTTTCAAGCTCTCAACGCCATCTACAGCTGCTTCCTTATCAGAAGCGCTT
                                                                                                                                                                 Conservative
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                                                                                                                                                                                31.7%;
87.7%;
                                                                                                                                                         Score 83.6; DB 17; Length 169;
Pred. No. 4.4e-17;
0; Mismatches 14; Indels 2
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Pred. No.
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1.1e-20;
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; OTHER INFORMATION: Clone ID: PAT_MRT3847_125713C.1
US-10-424-599-28474
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GENERAL INFORMATION:

APPLICANT: La Rosa Thomas J

APPLICANT: Kovalic David K

APPLICANT: Cao Yongwei

TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

FILE REFERENCE: 38-21(5323)B

FULE REFERENCE: 38-21(5323)B

CURRENT APPLICATION NUMBER: US/10/424,599

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 285684

SEQ ID NO 28474

LENGTH: 289
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US-10-424-599-86542
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                                                                                                                     Matches
                                                                                                                                   Query Match
Best Local (
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TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 86542
LENGTH: 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 28474, Application US/10424599 Publication No. US20040031072A1
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APPLICANT: Cao Yongwe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: La Rosa Thomas
APPLICANT: Kovalic David
                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Glycine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Clone ID: PAT_MRT3847_49159C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Glycine max FEATURE:
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75; Conserv
                                                                                                                                 Similarity
                                                                    TCTTAAAACCTCCTTACGGATCAAATGCTCCCAAACAGATGGGAATGGAAGTCCTGCAAA 241
GAGGACA-TGCTTCATGATCTTT
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                                                                                                                 Conservative
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                                                                                                             Score 58.2; DB 17;
Pred. No. 1.1e-08;
0; Mismatches 8;
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Pred. No. 1.9e-09;
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RESULT 13
US-10-021-323-1453
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                                                                                                                            PEATURE:

NAME/KEY: unsure

LOCATION: (1)..(603)

OTHER INFORMATION: unsure at all n locations

OTHER INFORMATION: Clone ID: LIB3825-003-Q1-K6-E10
US-10-021-323-1453
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TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 58838
LENGTH: 208
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Publication No. US20040123340A1
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                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                           APPLICANT: Feng, Paul C.C.
APPLICANT: Fincher, Karen L.
APPLICANT: Ziegler, Todd E.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21 (52274) B
CURRENT APPLICATION NUMBER: US/10/021,323
CURRENT FILING DATE: 2001-12-12
PRIOR APPLICATION NUMBER: US 60/255, 619
PRIOR FILING DATE: 2000-12-14
NUMBER OF SEQ ID NOS: 17880
                                             Matches
                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 1453
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Deikman
                                                                                                                                                                                                                                                           LENGTH: 603
TYPE: DNA
ORGANISM: Gossypium hirsutum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Clone ID: PAT_MRT3847_24142C.1
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Similarity 84.5%;
                                                                 Similarity
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TTCCCAATGGCTTCCGTTTCCAAGCTCTCAACGCCAAATCCACTTGCTTCCTTATCAGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGTTAAGACCGCCTTACGGATCGAGTGCTCCGAGACAGGTGGGAATGGAAGTCCTGCAGA 60
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                                                               18.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 52.8; DB 17;
pred. No. 5.9e-07;
0; Mismatches 12;
                                           Score 47.6; DB 18;
Pred. No. 4.4e-05;
0; Mismatches 104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 208;
                                             Indels
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RESULT 15
US-10-424-599-102259
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US-10-424-599-15721
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CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 15721
LENGTH: 183
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TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21 (5323)B CURRENT APPLICATION NUMBER: US/10/424,599 CURRENT FILING DATE: 2003-04-28 NUMBER OF SEQ ID NOS: 285684 SEQ ID NO 102259
                                                                                                                                                                                                                                           Sequence 102259, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION:
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Best Local
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                                                                                                                                                     APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
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ORGANISM: Glycine max
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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Match Length
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US-09-902-540-1273
US-09-9134-000C-2307
US-09-949-016-163223
US-09-949-016-163283
US-09-949-016-17791
US-09-949-016-17792
US-09-949-016-178915
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US-09-949-016-17888
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US-09-949-016-17888
US-09-949-016-17888
US-09-949-016-178911
US-08-956-1718-931
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200077,
17375, A
1177, Ap
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176915,
11868, A
14589, A
14589, A
17119, A
17119, A
1931, App
931, App
931, App
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15815, Ap
1273, Ap
2307, Ap
163223,
16318, A
16790, A
112881, A
11287, A
17272, A
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RESULT 2 US-09-902-540-5815 ; Sequence 5815, Application US/09902540 ; Patent No. 6833447 ; Patent No. 6833447 ; GENERAL INFORMATION: ; APPLICANT: Goldman, Barry S. ; APPLICANT: Hinkle, Gregory J. ; APPLICANT: Slater, Steven C. ; APPLICANT: Wiegand, Roger C. ; TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses The FILE REFERENCE: 38-10(15849)B	Query Match 30.3%; Score 91; DB 4; Length 269; Best Local Similarity 88.2%; Pred. No. 7.1e-22; Matches 105; Conservative 0; Mismatches 13; Indels 1; Gaps Qy 1 GATTTATTGACAACACCGATCCTGCTGGGATTGATCAAATTGCTCAACTAGGACCTG Qy 1 GATTTATTGACAACACAGATCCTGCTGGGATTGATCATCAAATTGCTCAACTAGGACCTG Db 124 GATTTATTGACAACACAGATCCTGCTGGGATTGATCATCAAATTGCTCAACTAGGACCTG Qy 61 AACTGGCAACTACTCTTGTAATTGTCATTTTCTAAGAGCGGAGCACCCTGAAACCCGCC Qy 1	ERIVED	28 30.8 10.3 579 3 US-09-631-547-1 Sequence 29 30.8 10.3 601 4 US-09-949-016-140812 Sequence 30 30.8 10.3 264358 4 US-09-949-016-15725 Sequence 20 31 30.6 10.2 1051 3 US-08-961-083-125 Sequence 31 30.6 10.2 1051 4 US-09-536-784-125 Sequence 31 30.6 10.2 2243 4 US-08-9561-787-33 Sequence 31 30.6 10.2 22243 4 US-08-956-1718-164 Sequence 31 30.6 10.2 22243 4 US-08-956-1718-164 Sequence 31 30.4 10.1 38239 4 US-09-949-016-12348 Sequence 31 30.4 10.1 38239 4 US-09-949-016-12348 Sequence 31 30.4 10.1 38252 4 US-09-949-016-12348 Sequence 31 30.2 10.1 39320 4 US-09-949-016-12461 Sequence 31 30.2 10.1 235064 US-09-949-016-16853 Sequence 31 30.2 10.1 235064 US-09-949-016-16853 Sequence 31 30 10.0 236815 4 US-09-949-016-16853 Sequence 31 30 10.0 24841 4 US-09-949-016-163182 Sequence 31 30 10.0 24841 4 US-09-949-016-17543 Sequence 31 30 10.0 36546
and	A 269; Lels 1; Gaps 1; GCTCAACTAGGACCTG 60 GCTCAACTAGGACCTG 183 ACACCTGAAACCCGC 119 ACCUNTAACCCCCC 241	ERIVED FRO	Sequence 1, Appli Sequence 140812, Sequence 15725, App Sequence 125, App Sequence 33, Appl Sequence 164, App Sequence 164, App Sequence 63183, A Sequence 12348, A Sequence 12461, A Sequence 12461, A Sequence 16853, A Sequence 16274, A Sequence 16274, A Sequence 63182, A Sequence 63182, A Sequence 16274, A Sequence 63182, A Sequence 63182, A Sequence 63182, A Sequence 63182, A Sequence 63182, A Sequence 17543, A

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; LOCATION: (1)..(72704)
; OTHER INFORMATION: unsure at all n locations
US-09-902-540-1273
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CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 5815
LENGTH: 1590
TYPE: DNA
ORGANISM: Myxococcus xanthus
US-09-902-540-5815
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Best Local Similarity 53.0%;
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APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
APPLICATI: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15849)B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 1273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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Best Local Similarity
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                                      CTGGCAACTACTCTTGTAATTGTCATTTCTAAGAGCGGAGGCACACCTGAAAACCCGGCAAT 122
                                                                                                                TTTATTGACAACACCGATCCTGCTGGGATTGATCATCAAATTGCTCAACTAGGACCTGAA
     CTGTCGGAGACGCTCACCCTGGTCATCAGCAAGTCGGGTGGCACCAAGGAGACGCGCAAC 33746
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53.0%;
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                                                                                                                                                                           Score 74;
Pred. No. 1
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Pred. No. 1.9e-15;
                                                                                                                                                    Mismatches 140; Indels
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1.5e-14;
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US-09-949-016-163223/c
; Sequence 163223, Application US/09949016
; Patent No. 6812339
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Best Local Similarity
Matches 63; Conserva
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                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION
FILE REFERENCE: CLOO1307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/241,756
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
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                                                          PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 163223
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PRIOR FILING DATE: 1997-08-15
NUMBER OF SEQ ID NOS: 6812
SOFTWARE: PatentIn version 3.1
SEQ ID NO 2307
LENGTH: 1200
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Patent No. 6617156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Lynn Doucette-Stamm et al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO TITLE OF INVENTION: ENTEROCOCCUS FAECALIS FOR DIAGNOSTICS AND THERAPP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/134,000C CURRENT FILING DATE: 1998-08-13
ORGANISM: Human
                    TYPE: DNA
                                         LENGTH:
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; NAME/KEY: misc_feature
; LOCATION: (1)...(80004)
; OTHER INFORMATION: n = A,T,C or G
US-09-949-016-16318
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GENERAL INFORMATION:
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                                                          CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
                                                                                                                                                                                                       APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
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PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                     PRIOR APPLICATION NUMBER: 60/231,498 PRIOR FILING DATE: 2000-09-08
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CURRENT FILING DATE: 2000-04-14
                                                                                                                                                                                        FILE REFERENCE: CL001307
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ilarity 56.4%;
Conservative
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Pred. No. 3.9;
0; Mismatches 48;
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Pred. No. 0.26;
0; Mismatches 48
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                                                                                                        US-09-949-016-14376
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
                                                            Sequence 14376, Ap
Patent No. 6812339
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; TYPE: DNA
; ORGANISM: Human
US-09-949-016-16790
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Best Local Similarity
Matches 60; Conserv
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SEQ ID NO 16790
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/231,498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2000-10-03
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                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (1)...(197336)
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 197336
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                                                                                                                                                                                                                         Local Similarity
les 69; Conserv
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No. 6812339
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                                   TAACACTGC
                                                                         TGTTGAGTGTACACTTAAAAATCAGGAGGTTTCACATGAAAATCTCAATTTTCAGTTCCT 89937
                                                                                                            TGCGGGGCTGCAATTCTCGAAACAGGGTGTTGCAATTACTCAAGAAAATTCTCTGTTGGA 214
                                                                                                                                                  GGGCGGCCTCGCCCCCCCCCCCAATGTTTTATTTGACTCGTACAACGGCTTCAGAAA
                                                                                                                                                                                                                             Conservative
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53.5%;
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Pred. No. 7.5;
0; Mismatches
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Pred. No. 4.5;
0; Mismatches
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OF DETECTION AND USES THEREOF
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Application US/09949016

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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THER
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIO
                                                                                                                                                                                                          ; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(234288)
; OTHER INFORMATION: n = A,T,C or
US-09-949-016-17272
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                                                                                                                                                       Query Match
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Best Local Similarity 53...
69; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 17272
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NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14376
LENGTH: 197337
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PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Human
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
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NAME/KEY: misc feature
LOCATION: (1)...(197337)
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95 GAGCGGAGGCACACCTGAAACCCCGCAATGGTCTACTAGAAGTACAGAAAGCCTTCAGAGA 154
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53.5%;
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                                                                         Score 33; DB 4; Length 234288; Pred. No. 8.2; 0; Mismatches 60; Indels 0
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US-09-949-016-176915/c
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                                   NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 11868
                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION FILE REFERENCE: CL001307
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 176915
                                                                                                                                                                                                                                                                                                                                                   Sequence 11868, Application US/09949016 Patent No. 6812339
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APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT PILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR PILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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                                                                                                                                               PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
                                                                                             PRIOR APPLICATION NUMBER: 60/231,498 PRIOR FILING DATE: 2000-09-08
                                                                                                                                    PRIOR FILING DATE: 2000-10-03
                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
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              ENGTH: 300598
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                                                   FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                   DETECTION
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US-09-949-016-14588
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; Sequence 14589, Application US/09949016
; Patent No. 6812339
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                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc feature
; LOCATION: (1) ... (302604)
; OTHER INFORMATION: n = A,T,C or G
US-09-949-016-14588
                                     US-09-949-016-14589
                                                           RESULT 14
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                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 14588
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PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FASTSEQ for Windows Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE: misc_feature
NAME/KEY: misc_feature
LOCATION: (1)...(300598)
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENGTH: 302604
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Similarity 52.6%;
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Pred. No. 13;
0; Mismatches
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Pred. No. 13;
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US-09-949-016-17119
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US-09-949-016-17119
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                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOPTWARE: FastSEQ for Windows Versio
SEQ ID NO 17119
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APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 14589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 6812339
                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
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NUMBER OF SEQ ID NOS: 207012
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PRIOR APPLICATION NUMBER: 60/231,498
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                NAME/KEY: misc_feature
LOCATION: (1)...(308362)
OTHER INFORMATION: n = A,T,C or G
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LOCATION: (1)...(302604)
OTHER INFORMATION: n = A,T,C
                                                                                                       ORGANISM: Human
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                                                                                                                                 TYPE: DNA
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71; Conserv
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llarity 52.6%;
Conservative
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                                                                                                                                                                                                    for Windows Version 4.0
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Pred. No. 13;
0; Mismatches
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OF DETECTION AND USES THEREOF
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Query Match Best Local Similarity

10.9%;

Score 32.6; Pred. No. 13

DB

Length 308362;

GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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length: 2000000000
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Gapop 10.0 , Gapext 1.0
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/cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US10F_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US11_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
                                                                                                                                                                                                                                                                                                                 6/ptodata/1/pubpna/USO8 NEW PUB.seq:*
6/ptodata/1/pubpna/USO8_PUBCOMB.seq:*
6/ptodata/1/pubpna/USO9A_PUBCOMB.seq:*
                                                                                                                                                                                                                                                                                                                                                                                    6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
-6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
-6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
-6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
                                                                                                                                                                                     6/ptodata/1/pubpna/US09C_PUBCOMB.seq:*
6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
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D_PUBCOMB.seq:*
E_PUBCOMB.seq:*
F_PUBCOMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.

Score

Query Match Length DB

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Description

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218.4	244	285.6 247.2	285.6	288.8	300
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US-10-767-795-1472	US-10-739-930-5223 US-10-437-963-92447	US-10-425-115-79836 US-10-437-963-92448	US-10-425-114-2691 US-10-425-114-35572	US-10-425-115-79832	US-09-237-183A-276
Sequence 1472, Ap	Sequence 5223, Ap Sequence 92447, A	Sequence 79836, A Sequence 92448, A	Sequence 2691, Ap Sequence 35572, A	Sequence 79832, A	Sequence 276, App

ALIGNMENTS

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CURRENT APPLICATION NUMBER: US/09/237,183A
CURRENT FILING DATE: 1999-01-26
PRIOR APPLICATION NUMBER: US 60/067,000
PRIOR FILING DATE: 1997-11-24
NUMBER OF SEQ ID NOS: 2814
SEQ ID NO 276
LENGTH: 300
TYPE: DNA
ORGANISM: Zea mays
US-09-237-183A-276
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                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 300; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Fisher, Dane K.
APPLICANT: Liu, Jingdong
TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With The
TITLE OF INVENTION: Sucrose Pathway
FILE REFERENCE: 38-21(15089)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Cheikh, No. US20030135870Aldine
                                61 AACTGGCAACTACTCTTGTAATTGTCATTTCTAAGAGCGGAGGCACACCTGAAACCCGGA 120
                                                                                                                           1 GATTTATTGACAACACCGATCCTGCGGGATTGATCATCAAATTGCTCAACTAGGACCTG
AACTGGCAACTACTCTTGTAATTGTCATTTCTAAGAGCGGAGGCACACCTGAAACCCGGA 120
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US-10-425-115-79832
; Sequence 79832, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 33220, Application US/10425114

Publication No. US20040034888A1

GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
ITILE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
ITILE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
SEQ ID NO 33220
LENGTH: 2175
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US-10-425-114-33220
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Best Local Similarity
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ORGANISM: Zea mays
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         768 GATTTATTGACAACACCGATCCTGCTGGGATTGATCATCAAATTGCTCAACTAGGACCTG 827
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Pred. No. 6.7e-88;
0; Mismatches 2
    Molecules
and Other Molecules Associated With
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APPLICANT: Liu, Jingdong
APPLICANT: APLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5313) B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 2691
LENGTH: 1822
TYPE: DNA
CDCANTEM: 702 marks
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; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 79832
; LENGTH: 2737
; TYPE: DNA
; ORGANISM: Zea mays
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; OTHER INFORMATION:
US-10-425-115-79832
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2691, Application US/10425114
Publication No. US20040034888A1
GENERAL INFORMATION:
                                                                                                                                            Matches 291;
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Best Local
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                                                                                                                                                                 Local Similarity
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                                                                               1 GATTTATTGACAACACCGATCCTGCTGGGATTGATCATCAAATTGCTCAACTAGGACCTG 60
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293; Conserv
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AACTGGCAACTACTCTTGTAATTGTCATTTCTAAGAGCGGAGGCACCACCTGAAACCCGCA
                                                           GATTTATTGACAACACAGATCCTGCTGGGATTGATCATCAAATTGCTCAACTAGGACCTG
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                                                                                                                                          Conservative
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97.0%;
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97.7%;
                                                                                                                                        Score 285.6; DB 17;
Pred. No. 3.3e-84;
0; Mismatches 9;
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Pred. No. 3.5e-85;
0; Mismatches 7;
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US-10-425-115-79836; Sequence 79836, Application US/10425115; Publication No. US20040214272A1; GENERAL INFORMATION:
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LENGTH: 3088
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APPLICANT: La Rosa, Thomas J. APPLICANT: Kovalic, David K. APPLICANT: Zhou, Yihua
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APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (5331)B
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
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ORGANISM: Zea mays
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                                                                                                                                                                                                                                                                                              GTGTTGCAATTACTCAAGAAAATTCTCTGTTGGATAACACTGCTAGAATAGAGGGATGGT 240
                                                                                                                                                                                                                                   TAGCTCGGTTTCCTATGTTTGATTGGGTTGGTGGTAGGACTTCAGAAATGTCTGCTGTGG 300
                                                                                                                                                                                                                                                                        GTGTTGCAATCACTCAAGAAAATTCTCTGTTGGATAACACTGCCAGAATAGAGGGATGGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95.2%;
97.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 285.6; DB 1
Pred. No. 4.3e-84;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 3088;
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                                                                                                                               US-10-437-963-92448
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CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 79836
LENGTH: 3830
TYPE: DNA
                                                        Query Match
Best Local Similarity
Matches 267; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 92448, Application US/10437963 Publication No. US20040123343A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 92448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 291;
                                                                                                                                                                                                                                                                      APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: La Rosa, Thomas J. APPLICANT: Kovalic, David K. APPLICANT: Zhou, Yihua
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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TITLE OF INVENTION: Nucleic Acid Molecules
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
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                                                                                                                                                                                                  TYPE: DNA
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                                                                                                                                            OTHER INFORMATION: Clone ID: PAT_MRT4530_90929C.1
                                                                                                                                                                                 ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION:
                                                                                                                                                               FEATURE:
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GATTTATTGACAACACCGATCCTGGCTGGGATTGATCATCAAATTGCTCAACTAGGACCTG
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Wu, Wei
Boukharov, Andrey
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Pred. No. 4.8e-84;
0; Mismatches 9
                                                          0;
                                                        Score 247.2; DB 18;
Pred. No. 2.2e-71;
0; Mismatches 33;
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Publication No. 1
GENERAL INFORMATION:
GENERAL INFO
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; LENGTH: 2696
; TYPE: DNA
; ORGANISM: Triticum aestivum
; FEATURE:
; OTHER INFORMATION: Clone ID: US-10-739-930-5223
                                                                                                                                                                                          RESULT 9
US-10-437-963-92447/c
; Sequence 92447, Application US/10437963
; Publication No. US20040123343A1
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US-10-739-930-5223
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APPLICANT: KOVALIC, DAVIG K.

TITLE OF INVENTION: NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED TITLE OF INVENTION: PLANTS AND USES THEREOF FOR PLANT IMPROVEMENT FILE REFERENCE: 38-21(53377)B

CURRENT APPLICATION NUMBER: US/10/739,930

CURRENT FILING DATE: 2003-12-18

NUMBER OF SEQ ID NOS: 11088
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Best Local
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88.3%;
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Pred. No. 3e-70;
0; Mismatches 35; Indels
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US-10-767-701-27794
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                                                                                                                                                                                                                                                               APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof For Plant Improvement
FILE REFERENCE: 38-21(53535)8
CURRENT APPLICATION NUMBER: US/10/767,701
CURRENT FILING DATE: 2004-01-29
NUMBER OF SEQ ID NOS: 63128
SEQ ID NO 27794
LENGTH: 423
TYPE- NA
                                                                                      Matches
                                                                                                          Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 27794, Application US/10767701
Publication No. US20040172684A1
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LENGTH: 2666
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                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Sorghum bicolor
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Boukharov, Andrey A.
APPLICANT: Barbazuk, Brad
APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
IITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
IITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Oryza sativa
FEATURE:
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hes 264;
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                     1 GATTTATTGACAACACCGATCCTGCTGGGATTGATCAACTAGCAACTTGCTCAACTAGGACCTG
                                                                                                        Similarity
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GATTTATTGACAACACCGATCCTGCTGGTATTGATCATCAAATTGCTCAACTAGGACCTG
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ilarity 88.0%;
Conservative
                                                                                                     75.1%;
98.7%;
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                                                                                Score 225.2; DB 1
Pred. No. 2.2e-64;
0; Mismatches 3
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                                                                                                                       DB 18;
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US-10-425-114-9307; Sequence 9307, Application US/10425114; Publication No. US20040034888A1
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US-10-767-795-1472
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APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof For Plant Improvement
FILE REFERENCE: 38-21(53534)B
CURRENT APPLICATION NUMBER: US/10/767,795
CURRENT FILING DATE: 2004-01-30
NUMBER OF SEQ ID NOS: 117596
SEQ ID NO 1472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1472, Application US/10767795
Publication No. US20040181830A1
GENERAL INFORMATION:
APPLICANT: Kovalic, David K.
                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
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Local Similarity 83.0%;
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APPLICANT: Kovalic, David K.

APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 29640
LENGTH: 2218
TYPE: NN"
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CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 9307
LENGTH: 1411
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                                                                                                                                       Matches 243;
                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                        ORGANISM: Glycine max
FEATURE:
OTHER INFORMATION: Clone ID: UC-GMFLMINSOY100E04_FLI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
OTHER INFORMATION: Clone ID:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Glycine max
                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    241
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GATTTATTGACAACACCGATCCTGCTGGGGATTGATCAACATTGCTCAACTAGGACCTG
                                                                                                                                                       Similarity
                                                                                      GATTTATTGACAACACGATCCTGCTGGGATTGATCATCAAATTGCTCAACTAGGACCTG
                              AACTGGCAACTACTCTTGTAATTGTCATTTCTAAGAGCGGAGGCACACCTGAAACCCGCA 120
                                                                  GATTTGTGGACAACACGGATCCTGCTGGAATTGATCATCAGATTGCACAACTTGGGCCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TAGCTAGATTTCCTATGTTTGATTGGGTGGGAGGTAGAACGTCAGAGATGTCTGCAGTTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATGGTCTACTAGAAGTACAGAAAGCCTTCAGAGATGCGGGGCTGCAATTCTCGAAACAGG 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AGCTAGCTTCAACACTTGTGATTGTGATATCAAAGAGTGGAGGTACTCCTGAGACCAGAA 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GATTTGTGGACAACACGGATCCTGCTGGAATTGATCATCAGATTGCACAACTTGGGCCTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GTGTTGCAATTACTCAAGAAATTCTCTGTTGGATAACACTGCTAGAATAGAGGGATGGT 240
                                                                                                                                         Conservative
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81.0%;
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                                                                                                                                    Score 208.8; DB 1
Pred. No. 1.5e-58;
0; Mismatches 57
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Pred. No. 1.2e-58;
0; Mismatches 57;
                                                                                                                                                                        DB 17; Length 2218;
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AGCTAGCTTCAACACTTGTGATTGTGATATCAAAGAGTGGAGGTACTCCTGAGACCAGAA

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RESULT 15
US-09-938-842A-1222
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B
Sequence 1222, Application US/09938842A Patent No. US20020160378A1
GENERAL INFORMATION:
APPLICANT: Harper, Jeff
APPLICANT: Kreps, Joel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)B
FULE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 11100
LENGTH: 2547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 11100, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION:
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Glycine max
PEATURE:
NAME/KEY: unsure
LOCATION: (1)..(2547)
OTHER INFORMATION: unsure at all n locations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                            1262
                                                                                                                                                                                                                                                                                                                                                                                                   1082 AGCTAGCTTCAACACTTGTGATTGTGATATCAAAGAGTGGAGGTACTCCTGAGACCAGAA
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                                                                                                                                                                  TAGCTAGATTTCCTATGTTTGATTGGGTGGGAGGTAGAACGTCAGAGATGTCTGCAGTTG 1321
                                                                                                                                                                                            TAGCTCGGTTTCCTATGTTTGATTGGGTTGGTGGTAGGACTTCAGAAATGTCTGCTGTGG
                                                                                                                                                                                                                                                                  GTGTTGCAATTACTCAAGAAAATTCTCTGTTGGATAACACTGCTAGAATAGAGGGATGGT 240
                                                                                                                                                                                                                                                                                                                                                 ATGGTCTACTAGAAGTACAGAAAGCCTTCAGAGATGCGGGGGCTGCAATTCTCGAAACAGG 180
                                                                                                                                                                                                                                                                                                                                                                                                                                 AACTGGCAACTACTCTTGTAATTGTCATTTCTAAGAGCGGAGGCACACCCTGAAACCCGCA 120
                                                                                                                                                                                                                                          GTGTTGCTÄTAÄCACAAGAAATTCTTTGTTGGATAACACTGCAAGAATTGAGGGCTGGT 1261
                                                                                                                                                                                                                                                                                                                       ATGGTTTATTGGAAGTGCAGAAGGCCTTTCGTGAAGCAGGCTTGGATTTTCCTAAACAGG 1201
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Pred. No. 1.6e-58;
0; Mismatches 57
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APPLICANT: Wang, Xun
APPLICANT: Zhu, Tong
ITITLE OF INVENTION: STRESS-REGULATED GENES OF PLANTS, TRANSGENIC PLANTS CONTAINING
ITITLE OF INVENTION: SAME, AND METHODS OF USE
ITITLE OF INVENTION: US/09/938,84ZA
CURRENT APPLICATION NUMBER: US/09/938,84ZA
CURRENT FILING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: US 60/227,866
PRIOR APPLICATION NUMBER: US 60/264,647
PRIOR APPLICATION NUMBER: US 60/264,647
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-06-22
INUMBER OF SEQ ID NOS: 5379
SEQ ID NO 1222
LENGTH: 1836
TYPE: DNA
COCANTSM: Arabidopsis thaliana
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Local Similarity 78.7%;
      188
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                                                                                                                                                                                                    121 ATGGTCTACTAGAAGTACAGAAAGCCTTCAGAGATGCGGGGCTGCAATTCTCGAAACAGG 180
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                                                                                                            GTGTTGCAATTACTCAAGAAAATTCTCTGTTGGATAACACTGCTAGAATAGAGGGATGGT 240
                                                                                                                                                                   ATGGACTACTGGAAGTGCAGAAAGCATTCCGTGAAGCTGGTCTGAATTTCGCAAAACAGG
                              TAGCTCGGTTTCCTATGTTTGATTGGGTTGGTGGTAGGACTTCAGAAATGTCTGCTGTGG 300
                                                                                                                                                                                                                                                                                          AACTGGCAACTACTCTTGTAATTGTCATTTCTAAGAGCGGAGGCACACCTGAAACCCGCA 120
                                                                                 GTGTTGCAATAACCCAAGAGAACTCATTACTGGATAATACGGCAAGAATTGAAGGTTGGC 880
TAGCTAGATTTCCTATGTACGACTGGGTGGGTGGAAGAACATCAATAATGTCTGCAGTTG 940
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Pred. No. 7.6e-55;
0; Mismatches 64;
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Search completed: May 4, Job time: 341.366 secs 2005, 21:11:52